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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
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                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                        Score
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seq length: 2000000000
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Match Length
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                                                                                                                                                                                                                                                                                                                                                                                 / cgn2_6/ptodata/2/pubpna/US07_NEW PUB. seq: *
/ cgn2_6/ptodata/2/pubpna/US07_NEW PUB. seq: *
/ cgn2_6/ptodata/2/pubpna/US08_NEW PUB. seq: *
/ cgn2_6/ptodata/2/pubpna/US08_NEW PUB. seq: *
/ cgn2_6/ptodata/2/pubpna/US08_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US09A_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US09B_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US09B_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US10A_PUBCOMB. seq: *
/ cgn2_6/ptodata/2/pubpna/US10E_PUBCOMB. seq: *
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/cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*
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US-10-365-623-16

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US-10-075-994A-3
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US-09-930-283A-3
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Sequence 15, Appl
Sequence 25, Appl
Sequence 16, Appl
Sequence 1, Appli
Sequence 3, Appli
                                                                                                                Sequence 1, Appli
Sequence 3, Appli
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-10-425-115- -10-424-599- -10-357-930-	US-09-732-627A-2120 US-10-437-963-4788 US-09-983-965-3423	-10-926-543-44 -10-936-273-30 -10-809-189-1135	-10-843-641A- -10-936-273-2	10-057 10-173 10-371	9-969-	-10-440-341-2	-10-767-795-	10-027-632- 10-027-632-	27-632- 27-632-	18-995- 29-386-	9-728-446-1067	US-10-809-189-113549	US-09-930-283A-2 US-10-075-994A-2	-10-173-225B-	0-925-734-1	-10-437-258-2	-10-437-275-2	US-10-437-924-1 US-10-437-263-25	-10-075-994A
Sequence 14903 Sequence 10304 Sequence 50238	Sequence 2120, Ap Sequence 4788, Ap Sequence 3423, Ap	30, Ap 113548	293	Sequence 25, Appl Sequence 64, Appli Sequence 1, Appli		e 361		244006	2440 2440		1067,	Sequence 113549,	Sequence 2, Appli Sequence 2, Appli	Ф	Sequence 15, Appl Sequence 49, Appl	o ·	25	Sequence 1, Appii Sequence 25, Appi	4.

ALIGNMENTS

NAME: Hendricks, Glenna REGISTRATION NUMBER: 32,535 REFERENCE/DOCKET NUMBER: kasid TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 591-4470

ATTORNEY/AGENT INFORMATION:

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (703) 591.4428
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                           NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: kasid
TELECOMVUNICATION INFORMATION:
TELEPHONE: (703) 591-4470
TELEPHONE: (703) 591-4428
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/930,283A
FILING DATE: 16-Aug-2001
CLASSIFICATION: -Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/354,109
FILING DATE: 1999-07-15
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 22031
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gokhale, Prafulla
Dritschilo, Anatoly
Rahman, Aqulur
TITLE OF INVENTION: Liposomes containg Oligonucelotides
NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
ANTI-SENSE: YES
                 TOPOLOGY: unknown MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kasid, Usha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 1:
HYPOTHETICAL: NO
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: VA
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Hendricks and Assoc. STREET: P.O. Box 2509 CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Pred. No.
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US-10-290-545-25

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; ANTI-SENSE: YES ; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-930-283A-3
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                                                                                                                                        ; SEQUENCE DESCRIPTION: SEQ ID NO: 15: US-09-895-480A-15
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Best Local Similarity 100.0%;
Matches 15; Conservative 0,
                                                                     Matches
                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Inex Pharmaceuticals Inc.
TITLE OF INVENTION: High Efficiency Encapsulation of Charged Therapeutic
                                                                                                                                                                                                                                                                                                    TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER TYPE: Diskette, 3.5 inch, 1.44 Mb
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/895,480A FILING DATE: 29-Jun-2001 CLASSIFICATION: «Unknown» PRIOR APPLICATION DATA:
                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: «Unknown»
REFERENCE/DOCKET NUMBER: «Unknown»
TELECOMMUNICATION INFORMATION:
TELEPHONE: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                           ANTI-SENSE: yes
                                                                                                                                                                                            HYPOTHETICAL: no
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                          1 GTGCTCCATTGATGC 15
1 GIGCICCATIGATGC 15
                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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ZIP: 80435
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STATE: CO
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STREET: PO Box 5068
                                                                                                                                                                                                                                                                                                                                                           TELEFAX: <Unknown>
                                                                                                                                                                                                                                                                                  LENGTH: 15
                                                                     Conservative
                                                                                   100.0%;
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Pred. No. 1.3e+02;
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Pred. No. 1.3e+02;
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                                                                                                      DB 10; Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: DNA; ORGANISM: Homo sapiens US-10-290-545-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-365-623-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16, Application US/10365623 Publication No. US20030166512A1 GENERAL INFORMATION:
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CURRENT FLILING DATE: 2002-11-07

NUMBER OF SEQ ID NOS: 30

SOFTWARE: Patentin version 3.1

SEQ ID NO 25

SEQ ID NO 25
                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                             Sequence 1, Application US/10075994A Publication No. US20030215489A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 16
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Xie, Dong
TITLE OF INVENTION: Protein Carrier System for Therapeutic Oligonucleotides
FILE REFERENCE: 63024.000001
TRE/IN/165.623
                                                                                                                 APPLICANT: PIE, Jin
APPLICANT: MEWANI, Rajshree
APPLICANT: AHMAD, Imran
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/365,623
CURRENT FILING DATE: 2003-02-13
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.1
FILE REFERENCE: 219604
CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
                                                        APPLICANT: DRITSCHILO, Anatoly
APPLICANT: RAHMAN, Aquilur
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Klimuk, Sandy APPLICANT: Yuan, Zuan-Ning
                                                                                                                                                                                  APPLICANT: GOKHALE, Prafulla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Semple, Sean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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                                                                                                                                                                                                       Usha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 15; DB 16; 100.0%; Pred. No. 1.3e+02;
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; OTHER INFORMATION: Synthetic US-10-075-994A-3
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SEQ ID NO 3
LENGTH: 15
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Best Local Similarity
Matches 15; Conserv
CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.2
SEQ ID NO 4
                                                                                                                                                                                                                             Sequence 4, Application US/10075994A
Publication No. US20030215489A1
                                                                                                                                                                                                                  GENERAL INFORMATION
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CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
                                                                            FILE REFERENCE: 219604
                                                                                         APPLICANT: DRITSCHILO, Anatoly
APPLICANT: RAHMAN, Aquilur
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
                                                                                                                                         APPLICANT:
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APPLICANT: GOKHALE, Prafulla
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SOFTWARE: PatentIn version 3.2
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15; Conserv
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AHMAD, Imran
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AHMAD, Imran
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ilarity 100.0%;
Conservative (
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100.0%; Pred. No.
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; LENGTH: 15
; TYPE: DNA
; ORGANISM: Artificial Sequence
; PEATURE;
; OTHER INFORMATION: Synthetic
US-10-075-994A-4
           APPLICANT: Tam, Ying K.
APPLICANT: Chikh, Ghania
APPLICANT: Chikh, Ghania
APPLICANT: Hope, Michael J.
TITLE OF INVENTION: PATHOGEN VACCINES AND METHODS FOR
FILE REPERENCE: A-72216/TAL
CURRENT APPLICATION NUMBER: US/10/437,263
CURRENT FILING DATE: 2003-05-12
PRIOR APPLICATION NUMBER: 60/379,343
PRIOR APPLICATION NUMBER: 60/379,343
PRIOR FILING DATE: 2003-04-04
PRIOR FILING DATE: 2003-04-04
PRIOR FILING DATE: 2003-04-04
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; Sequence 25, Application US/10437263
; Publication No. US20040009943A1
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CURRENT APPLICATION NUMBER: US/10/347,924

CURRENT FILING DATE: 2003-01-21

PRIOR APPLICATION NUMBER: US 09/354,109

PRIOR FILING DATE: 1999-07-15

PRIOR APPLICATION NUMBER: US 08/957,327

PRIOR FILING DATE: 1997-10-24

PRIOR APPLICATION NUMBER: US 60/041,192

PRIOR FILING DATE: 1997-03-21

NUMBER OF SEQ ID NOS: 1
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Publication No. US20030229040A1
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Best Local Similarity
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APPLICANT: Tam, Y:
PRIOR APPLICATION NUMBER: 60/454,298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Dristschilo, Anatoly
APPLICANT: Rahman, Aquilur
TITLE OF INVENTION: CATIONIC LIPOSOMAL DELIVERY SYSTEM AND THERAPEUTIC USE THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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Zhang, Chuanbo
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100.0%; Pred. No. 1.3e+02;
tive 0; Mismatches 0;
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Sequence 25, Application US/10437258

Publication No. US20040013649A1

GENERAL INFORMATION:

APPLICANT: Tam, Ying K.

APPLICANT: Tam, Ying K.

APPLICANT: Chikh, Ghania

APPLICANT: Chikh, Ghania

FILE OF INVENTION: CANCER VACCINES AND METHODS OF USING THE SAME

CURRENT APPLICATION NUMBER: US/10/437,258

CURRENT FILING DATE: 2003-05-12

PRIOR APPLICATION NUMBER: 60/40,646

PRIOR APPLICATION NUMBER: 60/460,646

PRIOR FILING DATE: 2003-04-04

PRIOR FILING DATE: 2003-04-04

PRIOR APPLICATION NUMBER: 60/454,298
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US-10-437-263-25
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APPLICANT: Semple, Sean
APPLICANT: Chikh, Sandra
APPLICANT: Chikh, Ghania
ITITLE OF INVENTION: WETHYLATED IMMUNOSTIMULATORY OLIGONUCLEOTIDES AND
ITITLE OF INVENTION: USING THE SAME
FILE REFERENCE: A-72158/TAL
CURRENT APPLICATION NUMBER: US/10/437,275
CURRENT FILING DATE: 2003-05-12
PRIOR APPLICATION NUMBER: 60/379,343
PRIOR FILING DATE: 2002-05-10
PRIOR APPLICATION NUMBER: 60/460,646
PRIOR APPLICATION NUMBER: 60/460,646
PRIOR APPLICATION NUMBER: 10/290,545
PRIOR APPLICATION NUMBER: 10/290,545
PRIOR APPLICATION NUMBER: 10/290,545
PRIOR APPLICATION NUMBER: 10/290,545
PRIOR FILING DATE: 2002-11-07
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US-10-437-275-25
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 25
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Best Local Similarity
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Best Local Similarity
Matches 15; Conserv
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US-10-925-734-15
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; ORGANISM: Homo sapiens
US-10-437-258-25
                                                                                                           US-10-925-734-15
                                    Matches
                                                   Query Match
Best Local Similarity
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LENGTH: 15
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NUMBER OF SEQ ID NOS: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Inex Pharmaceuticals Inc.
TITLE OF INVENTION: High Efficiency Encapsulation of Charged
                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 80435

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.5 inch, 1.4

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS 5.0

SOFTWARE: Word Perfect

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/925,734

FILING DATE: 24-Aug-2004

CLASSIFICATION: «Unknown»

PRIOR APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: <Unknown>
REFERENCE/DOCKET NUMBER: <Unknown>
TELECOMMUNICATION INFORMATION:
TELEPHONE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/895,480 FILING DATE: 29-Jun-2001 ATTORNEY/AGENT INFORMATION:
                                                                                                                        HYPOTHETICAL: no
ANTI-SENSE: yes
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: other nucleic acid
                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                  15; Conservative
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1 GTGCTCCATTGATGC 15
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                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Dillon
STATE: CO
                                                                                                                                                                                                                                                                                                                           TELEFAX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Oppedahl & Larson LLP
STREET: PO Box 5068
                                                                                                                                                                                                                                                    LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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Pred. No. 1.3e+02;
                                                 Score 15; DB 21;
Pred. No. 1.3e+02;
                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.5 inch, 1.44 Mb
                                                                  Length 15;
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RESULT 15
US-10-173-225B-47
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NUMBER OF SEQ ID NOS: 109
SEQ ID NO 47
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 47, Application US/10173225B Publication No. US20030119769A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 49
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local
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                                                                                                            PRIOR FILING DATE: 1996-11-26
PRIOR APPLICATION NUMBER: PCT
PRIOR FILING DATE: 1995-05-31
                                                                                                                               PRIOR APPLICATION NUMBER: US 08/888,982
PRIOR TILING DATE: 1997-07-07
PRIOR APPLICATION NUMBER: US 08/756,806
PRIOR FILING DATE: 1996-11-26
PRIOR APPLICATION NUMBER: PCT/US95/07111
                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: US 10/057,550
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: US 09/143,214
PRIOR APPLICATION NUMBER: PCT/US98/13961
PRIOR APPLICATION NUMBER: PCT/US98/13961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/057,550
CURRENT FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: 09/506,073
PRIOR FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: US 09/143,214
PRIOR FILING DATE: 1998-08-28
                                                                     PRIOR APPLICATION NUMBER: US 08/250,856
PRIOR FILING DATE: 1994-05-31
                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1998-07-06
                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Antisense Oligonucleotide Modulation FILE REFERENCE: ISPH-0665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US98/13961
PRIOR FILING DATE: 1998-07-06
PRIOR APPLICATION NUMBER: US 08/888,982
PRIOR FILING DATE: 1997-07-07
PRIOR APPLICATION NUMBER: US 08/756,806
PRIOR FILING DATE: 1996-11-26
PRIOR APPLICATION NUMBER: PCT/US95/07111
                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/173,225B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Monia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 08/250,856 PRIOR FILING DATE: 1994-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Antisense Oligonucleotide Modulation of raf Gene Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1995-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: artificial sequence
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15; Conserv
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Pred. No. 1.3e+02;
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                 Database
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                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seq length: 0 seq length: 50
                                                                                                                                                                                                                                                                                                                               N Geneseq 16Dec04:*
1: geneseqn1980s:*
2: geneseqn1990s:*
3: geneseqn2000s:*
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SUMMARIES
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207.953 Million cell updates/sec
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15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	Score
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20	20	20	20	20	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	Length
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ADF09751	AAD44760	AAA73535	AAZ11557	AAT27527	ADR88950	ADL70154	ADF42926	ADF32025	ADE39690	ADE90171	ADF82830	ADB97456	ADB97458	ADA24233	ACC58517	AAD22797	AAZ98661	AAV99435	AAV54043	ID
Adf09751 Mouse/rat	Aad44760 Mouse and	Aaa73535 Mouse and	Aaz11557 Mouse and	Aat27527 Mouse/rat	Adr88950 Anti c-ra	Adl70154 Oligonucl	Adf42926 Methylate	Adf32025 Antisense	Ade39690 Oligonucl	Ade90171 Human c-r	Adf82830 Immunosti	Adb97456 Antisense	Adb97458 Sense (AT	Ada24233 Human c-r	Acc58517 Oligonucl	Aad22797 Human c-r	Aaz98661 Human c-r	Aav99435 Antisense	Aav54043 Human ant	Description

New cationic liposome composition containing raf oligodeoxynucleotide can be used to directly target tumour tissue and is useful in the radiation therapy of cancers.

WPI; 1998-532155/45.

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44.1	44	43	42	41	40	39	38	37	36	35	34	ω G	32	31	30	29	28	27	26	25	24	23	22	21
12	12	12	12	12	12	12	12.4	12.4	12.4	12.4	12.4	12.4	12.4	13	13	13	13	13	13	13	13	13	15	15
80.0	80.0	80.0	80.0	80.0	80.0	80.0	82.7	82.7	82.7	82.7	82.7	82.7	82.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	86.7	100.0	100.0
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AAD44716	AAA73491	AAA92026	AAA92027	AAZ11513	AAT86617	AAT27483	AAV93971	ABI85033	ABI85032	ABI94231	AAV23613	AAX16583	AAX16555	ACD42073	ADF09706	AAD44715	AAA73490	AAZ11512	AAT62145	AAT59716	AAT27482	AAV90935	ADB97457	ACD42120
Human	Aaa73491 Human c-r	Aaa92026 C-raf tar	Aaa92027 C-raf tar	Aaz11513 Human c-r	Aat86617 Rat c-raf	Aat27483 Human c-r	Aav93971 Human IL-	Abi85033 Capture o	Abi85032 Capture o	Abi94231 Capture o	Aav23613 Homo sapi	Aax16583 Position	Aax16555 Position	Acd42073 Antisense	Adf09706 Human c-r	Aad44715 Human c-r	Aaa73490 Human c-r	Aaz11512 Human c-r	Aat62145 Human c-r	Aat59716 Human raf	Aat27482 Human c-r	Aav90935 Human C-r	Adb97457 Oligo use	Acd42120 Antisense

ALIGNMENTS

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RESULT 1
AAV54043
21-MAR-1997;
24-OCT-1997;
                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                     Human; antisense; c-raf-1; oligodeoxyribonucleotide; ODN/oligo; tumour tissue; cancer; radiation therapy; radiosensitise; antisense;
                                                      Kasid U,
                                                                                                             19-MAR-1998;
                                                                                                                              01-OCT-1998.
                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                               Human antisense c-raf-1 oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                              02-DEC-1998 (first entry)
                                                                                                                                                                                                                                                                                                               AAV54043;
                                                                                                                                                                                                                                                                                                                             AAV54043 standard; DNA; 15
                                                                                                                                             W09843095-A1
                                                                                                                                                                                                                                              liposome carrier system; ss.
                                                                     (GEOU ) UNIV GEORGETOWN.
                                                      Gokhale P,
                                                                                      97US-0041192P.
97US-00957327.
                                                                                                              98WO-US005303
                                                                                                                                                             /*tag=
/note=
                                                                                                                                                                                                             Location/Qualifiers
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/note= "N-terminal base is phosphothioated"
15
                                                       Dritschilo
                                                                                                                                                             "C-terminal base is phosphothioated"
                                                                                                                                                                                                                                                                                                                              ВP
                                                      Þ
                                                       Rahman A;
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Claim

4

Page 21;

25pp;

English

This is the nucleotide sequence of the human antisense c-raf-1 oligodeoxyribonucleotide (ODN/oligo), used in the method of the invention to directly target tumour tissue, and in cancer radiation therapy. The products can be used in a method of radiosensitising tumour tissue by addition of an antisense oligonucleotide of maximum 40 bases containing ODN/oligo. The liposome carrier system directly targets tumour tissue and has the potential for use in the radiation therapy of cancers

밁

GTGCTCCATTGATGC GTGCTCCATTGATGC

15 15

AAV99435;

22-MAR-1999

(first

entry

aberrant gene

phosphorothioate;

Antisense oligonucleotide; human c-raf-1 protein kinase

expression; treatment; inflammation; infection; ss.

phosphodiester; lipid-encapsulation;

gene;

gene

Antisense oligonucleotide directed against c-raf-1 protein kinase

AAV99435 standard;

DNA;

15

ВP

Query Match Best Local S Matches 15

Similarity

100.0%;

Score 15; Pred. No.

15; μ

Conservative

0,

Mismatches

0

Gaps

0

1.le+02;

DB 2;

Length 15 Indels

Sequence 15

BP; 2 A; 4 C; 4 G; 5 T; 0 U;

0 Other;

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RESULT 2
ANV99435
ID ANV9
XX ANV9
XX ANV9
XX ANT1
DE ANT1
XX WO91
XX WO91
XX WO91
XX I14-1
XX IIII
XX II
                 The present sequence represents an antisense oligonucleotide directed against the human c-raf-1 protein kinase gene. The oligonucleotide can have either phosphorothioate or phosphodiester bonds. The oligonucleotide is lipid-encapsulated using the method of the invention. A composition comprising lipid-encapsulated particles of a therapeutic agent, e.g. antisense oligonucleotides, is prepared by mixing at least 2 lipids with buffered aqueous solution of charged therapeutic agent to form an intermediate mixture of lipid-encapsulated particles, and changing the pH of the mixture to neutralise at least some of the external surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Semple SC,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 23; 98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     Composition containing lipid-encapsulated therapeutic agent - use e.g. for delivering antisense molecules or ribozymes or treating associated with aberrant gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-045179/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-00856374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-CA000485
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Harasym
                                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 or phosphodiester bonds"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ansell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cullis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   useful,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    diseases
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This sequence represents an antisense oligonucleotide human c-raf-l protein kinase as its target gene. The c

e sequence which oligonucleotide

hae is

Example 5; Page 57; capsules comprising

99pp; English.

for delivering

ring antisense oligonucleotides steric barrier lipids.

ç

cells

using lipid

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RESULT 3
ANZ98661
ID ANZ9
XX ANZ9
XX ANZ9
XX Huma
XX Huma
XX Anti
KW Luma
XX Huma
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XX I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        charges on the particles. One lipid has a (de)protonatable group with Ka such that the lipid is charged at a first pH but neutral at a second pH (particularly near physiological pH) and the buffer maintains this lipid in the charged form (i.e. cationic when the therapeutic agent is anionic in the buffer, or vice versa). The second lipid prevents particle aggregation during formation of the lipid-therapeutic agent particles. The composition is used to introduce therapeutic agents into cells, in vivo or in vitro, particularly to treat or prevent diseases associated with aberrant gene expression in mammals, specifically tumours,
                                                                                                                                                                                                                                                     Klimuk SK,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CA2271582-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; phosphorothioate; inflammatory disease; tumour; gene therapy; aberrant gene expression; treatment; infectious disease; protein kinase C alpha; c-raf-1 protein kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15 BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                   WPI; 2000-225058/20.
                                                                                                                                                                                                                                                                                                                                                                          (ANSE/)
(CULL/)
(MOKW/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-NOV-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ98661 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation or infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HOPE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HARA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (KLIM/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              c-raf-1
                                                                                                                                                                                                                                                                                                                         HOPE M J.
ANSELL S M.
CULLIS P R.
MOK W W K.
SCHERRER P.
SEMPLE S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        \mathbf{L}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GTGCTCCATTGATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       KLIMUK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HARASYM T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                                  Harasym
Semple
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PK therapeutic antisense oligonucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-00078955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99CA-02271582
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Optionally phosphorothicate
                                                                                                                                                                                                                                                     SC,
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                                                                                                                                                                                                                                                                            Hope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15; DB 2;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                            Ansell
                                                                                                                                                                                                                                                                            MS,
                                                                                                                                                                                                                                                                               Cullis PR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein kinase;
                                                                                                                                                                                                                                                                                 MOX
MOX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 4
AAD22797
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The invention relates to a composition useful for treatment of e.g. tumours. The composition comprises lipid-therapeutic agent particles comprising a lipid portion and a charged therapeutic agent which is encapsulated in the lipid portion. The lipid portion comprises a first lipid component selected from lipids containing a protonatable or deprotonatable (preferably protonatable) group that has a pKa such that the lipid is in charged form at a first pH and in neutral form at a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 used in a method for delivering lipid encapsulated therapeutic agents (i.e antisense oligonucleotides) to mammals. The lipid capsule comprises steric barrier lipids that prevent particle aggregation during lipid nucleic acid formation. The method may be used for the delivery of therapeutic agents to mammalian cells. It is especially suitable for delivering nucleic acid molecules, and in particular antisense molecules which may be administered to down regulate the expression of aberrant genes. The aberrant gene may be ICAM-1, c-myc, c-myb, ras, raf, erb-B-2, PKC-alpha, IGF-IR, EGFR, VEGF and/or VEG-R-1. The method may be used for
                                                                                                                                                                                                        Disclosure; Col 15-16;
                                                                                                                                                                                                                                                       comprising lipid portion and a
                                                                                                                                                                                                                                                                               Composition useful for treatment of e
                                                                                                                                                                                                                                                                                                                                 WPI; 2002-024658/03.
                                                                                                                                                                                                                                                                                                                                                                                   Semple SC,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treatment; tumour; lipid-therapeutic agent particle; sphingomyelin; distearoylphosphatidylcholine; palmitoylcheoyl phophatidylcholine; DSPC; POPC; 1.2-diolecyl.sn.-phosphoethanolamine; cholesterol; SM; DOPE; inflammation; c-raf-1 protein kinase gene; human; infectious disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US6287591-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INEX-) INEX PHARM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human c-raf-1 protein kinase antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD22797;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD22797 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of tumours, inflammatory diseases and/or infectious
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ب
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                                                                                                                                                                 Klimuk SK,
Debeyer D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-00856374.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-00078954
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                   48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         Harasym T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Optionally phosphorothicate backbone"
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Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                     charged
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                                                                                                                                                                                                                                                       therapeutic
                                                                                                                                                                                                                                                                             tumors
                                                                                                                                                                                                                                                                                                                                                                                                           Ansell SM,
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                                                                                                                                                                                                                                                     comprises
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     pKa such that form at a
                                                                                                                                                                                                                                                                          particles
                                                                                                                                                                                                                                                                                                                                                                                                           Cullis
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RESULT 5
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ID ACCS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    during lipid therapeutic agent particles formation and which exchange out the lipid particle at a rate greater than PEG-CerC20; third lipid component is a neutral lipid selected from distearcylphosphatidylcholine (DSPC), palmitoylolecyl phophatidylcholine (POPC), 1,2-diolecyl-sn-3-phosphoethanolamine (DOPE) or SM (sphingomyelin) and a fourth lipid component which is cholesterol. Compositions of the invention are used for treatment or prevention of a disease caused by aberrant expression of a gene preferably ICAM-1 (Intracellular adhesion molecule-1), c-myc, c-myb, ras, raf, erb-B-2, PKC-alpha (phosphokinase C-alpha), IGF-IR (insulin growth factor 1-receptor), bcl-2, EGFR (epidermal growth factor receptor), VGGF and VGGF-R-1 (vascular endothelial growth factor receptor)), in a mammal or by inflammations such as tumour or an infectious disease. The present sequence is an antisense oligonucleotide targetted to human c-raf-1 protein kinase gene
                                                                                                        Improved mucosal adjuvant useful in the preparation of vaccine stimulating an immune response comprises a lipid-nucleic acid containing a nucleic acid component encapsulated by a lipid.
                                                                                                                                                                                                                                                                                                                                                                                                     07-NOV-2001; 2001US-0337522P.
10-MAY-2002; 2002US-0379343P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lipid nucleic
C-Raf-s; ss.
                                                           Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-NOV-2002; 2002WO-CA001717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide ODN #25 (hC-Raf1).
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                                                                                                                                                                                                                                WPI; 2003-493235/46.
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                                                                                                                                                                                                                                                                                      Klimuk S,
                                                        Page 21; 71pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 C; 4 G;
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Pred. No.
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The present sequence is that of oligodeoxynucleotide ODN #25 (hC-Raf-1)

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Best Local S
Matches 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for human C-Raf-s. It is an example of an ODN that can be used in lipid-nucleic acid (LNA) formulations of the invention comprising a lipid component and a nucleic acid component. The invention is based on the discovery that such LNA formulations associated with a target antigen stimulate enhanced mucosal immune responses, especially IgA production, directed to that target antigen in vivo as compared to the target antigen alone or mixed with free or unencapsulated forms of the ODN. Claimed improved mucosal vaccines comprise an LNA formulation with at least one antigen, the LNA formulation comprising a lipid component that encapsulates the nucleic acid component, with the lipid and nucleic acid components acting synergistically to stimulate antigen-specific IgG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapeutic oligonucleotide; double-stranded RNA; dsRNA; mobile protein; cytostatic; immunosuppressive; virucide; anti-HIV; antibacterial; cardiant; hyperproliferation; cancer; haematological; metastatic; autoimmune disease; infection; endocrine; neural; cardiovascular; pulmonary; reproductive system disorder; endocytosis; metabolic process; murine; intracellular adhesion molecule 1; ICAM-1;
                                                                                  Treating diseases with oligonucleotides or interfering RNA, useful for cancer or autoimmune diseases, covalently coupled to mobile pro in vivo or in vitro.
                                                                                                                                              WPI; 2003-646491/61.
                                                                                                                                                                             Xie D;
                                                                                                                                                                                                                                       13-FEB-2002; 2002US-0356053P
                                                                                                                                                                                                                                                                     13-FEB-2003; 2003WO-US004323
                                                                                                                                                                                                                                                                                                       21-AUG-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antisense oligonucleotide; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human c-raf-1 protein kinase antisense oligonucleotide
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The present invention describes a method for treating administering: (a) a therapeutic oligonucleotide (TON)

a disease by
) or double-stranded

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chemotherapeutic

Claim 128; Page 12; 42pp; English

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ADB97458/c
ID ADB974
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AC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 7
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   Chemosensitization of leukemia, lymphoma or
                                                                                                                WPI; 2003-689738/65.
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                                                                                                                                                                                                                                                                                                                                                                                                                   15-FEB-2002; 2002US-00075994
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                                                                                                                                                                                                                                Gokhale P,
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   myeloma, comprises administering a
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Pred. No. 1.1e+02;
                                                                                                                                                                                                                                Mewani R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local !
                                                                                                                                                                                                                                                                                                                                                                dimyristoyl trimethyl ammonium propane; DMTAP; phosphatidylcholine; PC cholesterol; cancer; leukaemia; lymphoma; myeloma; carcinoma; sarcoma; combination therapy; pre-cancerous lesion; chemotherapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                               antisense; ATG-AS; raf ODN; chemosensitisation; tumour tissue; chemotherapeutic agent; cationic liposome; cationic lipid; phosphatidylcholine; cholesterol; liposome; dimethyldioctadecyl ammonium bromide; DDAB;
                           Chemosensitization of tumor tissue, useful for treating leukemia, lymphoma or myeloma, comprises administering agent and cationic liposomes containing oligonucleotides {\bf r}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB97456 standard; DNA; 15
Example 1; Page 18; 77pp; English
                                                                                            WPI; 2003-689738/65.
                                                                                                                         Rahman
                                                                                                                                                                                                               15-FEB-2002; 2002US-00075994
                                                                                                                                                                                                                                          14-FEB-2003; 2003WO-US004681
                                                                                                                                                                                                                                                                           28-AUG-2003
                                                                                                                                                                                                                                                                                                         WO2003070221-A1
                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 15
                                                                                                                                       Kasid U,
                                                                                                                                                                  (GEOU )
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15; Conserv
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                                                                                                                                     Gokhale P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ATG-AS) raf ODN oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP;
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                                                                                                                                     Pei J,
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Pred. No.
                                                                                                                                     Mewani R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 T;
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thes 0;
                              oligonucleotides
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                                                            cancer,
                                            chemotherapeutic
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RESULT 9
ADF82830
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Best Local Similarity
                                         New pathogen vaccine having a lipid-nucleic acid formulation in combination with at least one microbial antigen, useful for stienhanced responses against bacterial, viral and parasitic infec
                                                                                                                                                                                            10-MAY-2002; 2002US-0379343P
07-NOV-2002; 2002US-00290545
12-MAR-2003; 2003US-0454298P
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                                                                                                                                  Semple
                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C-Raf-8;
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                                                                                                      WPI; 2003-903935/82
                                                                                                                                                                                                                                                         12-MAY-2003; 2003WO-CA000680
                                                                                                                                                                                                                                                                                                                     WO2003094829-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulant ODN25, component of lipid-nucleic acid vaccine
                                                                                                                                                                 (INEX-) INEX PHARM CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                  note= "OTHER= optional phosphorothioate nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                  /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vaccine; lipid-nucleic acid; phosphorothioate; human;
                                                                                                                                  Норе
                                                                                                                                                                                                                                                                                                                                                                   base= OTHER
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25;

138pp; English

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RESULT 10
ADE90171
ID ADE90
X S X P P P P X D X X D X S C X R R X R R X S C X P R X S C X P R X S C X P R X S C X P R X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C X S C 
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; lipid-encapsulated therapeutic agent particle; aberrant gene expression; intercellular adhesion molecule; ICAM-1; c-myc; c-myb; ras; raf; erb-B-2; protein kinase C; PKC-alpha; insulin-like growth factor; IGF-IR; epidermal growth factor receptor; EGFR; vascular endothelial growth factor; VEGF; VEGF-R-1; tumour;
                                                                            Preparation of a composition comprising lipid-encapsulated therapeutic agent particles, useful for introducing a nucleic acid into a cell and for treating diseases characterized by aberrant gene expression
                                                                                                                                                                                                                                               Semple SC,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KLIM/)
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                                                                                                                                                                                                                                                                                                                                                          (CULL/)
                                                                                                                                                                                                                                                                                                                                                                                                                (HOPE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HARA/)
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HOPE M J.
ANSELL S M.
CULLIS P.
SCHERRER P.
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KLIMUK S K.
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Debeyer D;
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98US-00078954.
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Pred. No.
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Disclosure; SEQ

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15;

52pp; English

The present invention describes a cancer vaccine (I), which comprises a lipid-nucleic acid (LNA) formulation in combination with at least one tumour-associated antigen that is mixed with or associated with the LNA formulation comprising a lipid component having at least one cationic lipid, and a nucleic acid component comprising at least one

origonucleotide,

where the vaccine is capable

of.

a Th-1 based

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ADE3 9690

ADE3 9690

AC ADE3

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Best Local S
Matches 15
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                                                                                                                                                                                                                                                              New cancer vaccine having a lipid-nucleic acid formulation in combination with at least one tumor-associated antigen, useful for stimulating
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07-NOV-2002; 2002US-00290545
04-APR-2003; 2003US-0460646P
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Pred. No. 1.1e+02;
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                                                                                                                                                                                    English
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                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to reducing the platelet count in a patis comprises preparing a formulation of an oligonucleotide with an agent that enhances penetration of the oligonucleotide into cells, and administering the formulation to a patient having an elevated platelet count. The oligonucleotide is useful for preparing a medicament for reducing the platelet count in a patient, particularly for treating the phatelet count in a patient, particularly for treating thrombocytosis. The present sequence represents an oligonucleotide of the plate of the plat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immune response in vivo to the at least one tumour-associated anti (I) has cytostatic activity, and can be used in vaccines, and in 9 therapy. The methods and compositions of the present invention can used for stimulating enhanced responses against tumour-associated antigens and for inhibiting tumour growth. The present sequence represents an oligonucleotide which is used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reducing the platelet count in a patient, useful for treating thrombocytosis, comprises administering antisense oligonucleotides inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with an agent that enhances penetration of the inhibiting raf-1 gene with a penetration of the inhibiting raf-1 generation of the i
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                                                                                                                                                                                                                                         stimulating an enhanced host immune response to antigenic stimulation, comprising administering to the host the LNA formulation; (4) stimulating host dendritic cells in vivo, comprising contacting at least one dendritic cells with the lipid-methylated nucleic acid formulation to a host; and (5) simultaneously delivering antigenic and adjuvant immune stimulation to antigen presenting cells, comprising the administration of the LNA formulation associated with a target antigen. The lipid-methylation nucleic acid formulation has immunostimulant, cytostatic, antiinflammatory and antiarthritic activities, and can be used in vaccines, and in gene therapy. The formulation and methods are useful in stimulating and/or expanding dendritic cell populations in response to antigenic stimulation. They may be used for treating cancer, inflammation, arthritis or immunodeficiency disorders. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            formulation for stimulating an immune response in an animal, compri-
lipid component and a nucleic acid component which is a methylated
nucleic acid sequence. Also described: (1) an adjuvant comprising
-nucleic acid (LNA) formulation; (2) a vaccine comprising the LNA
formulation in combination with at least one target antigen; (3)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lipid-methylated nucleic acid formulation for stimulating an immune response in an animal comprises a lipid component and a nucleic acid component comprising a methylated nucleic acid sequence.
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07-NOV-2002; 2002US-00290545.
04-APR-2003; 2003US-0460646P.
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arthritis; immunodeficiency disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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                                                                                                                          Sequence 15
                                                                                                                                                                            sequence represents a methylated immunostimulatory oligonucleotide given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 25; 102pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lipid-methylated nucleic acid formulation; immune response; lipid-nucleic acid; vaccine; immunostimulant; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INEX-) INEX PHARM CORP.
15;
                              Similarity
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                                                                                                                          BP; 2 A; 4 C; 4 G; 5 T; 0 U;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a lipid-methylated nucleic acid
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                           100.0%;
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                           Score 15; DB 12;
Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                              invention relates to novel gemcitabine compositions and their use in treating proliferative diseases such as cancer, particularly in mammals especially in humans. The compositions include liposome-entrapped gemcitabine. The cancer is especially lymphoma, ovarian cancer, breast cancer, pancreatic cancer, lung cancer or colon cancer. The liposomal gemcitabine compositions can be used in conjunction with secondary therapeutic agents including antineoplastic, antifungal and antibiotic agents as well as antisense oligonucleotides, especially an antisense oligonucleotides, especially an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treatment of cellular proliferative disease e.g. cancer involves administration of a composition comprising liposomal gemcitabine negatively charged phospholipid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide antisense to raf
     WO2004071466-A2
                                Synthetic.
                                                        C-raf-1; liposomal; antineoplastic; cytostatic; cancer; antisense; ss.
                                                                                    Anti c-raf-1 oligonucleotide
                                                                                                               18-NOV-2004
                                                                                                                                          ADR88950;
                                                                                                                                                                   ADR88950 standard;
                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                       oligonucleotide to raf (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of an antisense oligonucleotide to raf. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; SEQ ID NO 1; 25pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zhang J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-AUG-2002; 2002US-0405378P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Raf; antisense; liposome; drug delivery; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAY-2004
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                                                                                                                                                                                                                                                                                                        Local Similarity
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                                                                                                                                                                                                                                                     GTGCTCCATTGATGC 15
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                                                                                                                                                                                                                                                                                          Conservative
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Pred. No. 1.1e+02;
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                                                                                                                                                                                    composition to obtain a size-reduced liposomal composition, freeing the liposomal composition of the water-miscible organic solvent by disfiltration using a tangential flow filtration process and sterile filtration, sterile-filtering the liposomal composition and freeze-drying the liposomal composition and freeze-drying the liposomal composition and freeze-drying the liposomal preparation. Step (b) involves adding the water-miscible organic solvent solution to the aqueous solution while mixing and mixing of solution following addition of water-miscible solvent comprising the lipid fraction to the aqueous solution while cooling. The active principal comprises at least one antineoplastic or antifungal agent (preferably traxane, camptothecin or their derivatives, especially paclitaxel or docetaxel). The composition is used for the treatment of disease e.g. cancer. The composition eliminates the disease or its symptoms, need not completely eradicate the effects of the disease, reduces the severity of a disease, infection or reduction in the rate by which a disease progresses within a patient. The method permits the production of liposomal formulation on a commercial scale. The present sequence represents an antisense oligonucleotide specific for c-raf-1, that can be used as an active principal.
                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         method involves: dissolving a lipid fraction in a water-miscible organic solvent; and mixing the water-miscible organic solvent solution comprising the lipid fraction with an aqueous solution under conditions to form a bulk liposomal composition. The method further involves adding at least one active principal to the water-miscible organic solvent prior to the addition of the lipid fraction, or to the aqueous solution prior, during or after the step (b), size-reducing the bulk liposomal
                                                                                                                                                                                    Sequence 15 BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-FEB-2003; 2003US-0446895P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to the preparation of a liposomal composition. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 6; 27pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Preparation of liposomal composition used for treating e.g. cancer involves dissolving lipid fraction in water miscible organic solven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-AUG-2004.
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                                                                                                                  Similarity
                                         GTGCTCCATTGATGC 15
GTGCTCCATTGATGC 15
                                                                                          Conservative
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Copyright (c) 1993 - 2005 Compugen Ltd.
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11.8	11.8	11.8	11.8	11.8	11.8	11.8	11.8	11.8	11.8	12	12	12	12	12	12	12	12.4	12.4	12.4	12.4	12.4	12.4	12.4	13	13
78.7	7.8.7	78.7	78.7	78.7	78.7	78.7	78.7	78.7	78.7	80.0	80.0	•	80.0	80.0	80.0	80.0	82.7	82.7	82.7	82.7	82.7	82.7	82.7	86.7	86.7
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BD014421 Method fo	AR069912 Sequence	AR148361 Sequence	AR123544 Sequence	AR122889 Sequence	AR092534 Sequence	AR080410 Sequence	A16267 Oligonucleo		A16281 Oligonucleo	AR493387 Sequence	AR215956 Sequence	I27233 Sequence 3	E49513 Antisense o	AR106991 Sequence	AR106990 Sequence	AR073934 Sequence	CQ749113 Sequence	AR039324 Sequence	AX289556 Sequence	AX598466 Sequence	AX294189 Sequence	AR070338 Sequence	AR037100 Sequence	AR215955 Sequence	127232 Sequence 2

ALIGNMENTS

AR110775 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE REFERENCE AUTHORS RESULT 2 AR110777/c LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM JOURNAL FEATURES 밁 S ORIGIN Matches Query Match Best Local (source AR110777 Sequence 3 AR110777 AR110777.1 Unknown. Unclassified. 1 (bases 1 to 15) 1 (kasid,U., Gokhale,P., Dritschilo,A. and Liposomes containing oligonucleotides I (bases 1 to 15) Kasid, U., Gokhale, P., Dritschilo, A. and Rahman, A. Liposomes containing oligonucleotides Patent: US 6126965-A 1 03-OCT-2000; Location/Qualifiers 1 GTGCTCCATTGATGC Similarity Unknown Unclassified. Unknown Unknown. AR110775.1 Sequence 1 from patent US AR110775 AR110775 GTGCTCCATTGATGC 100.0%; ilarity 100.0%; Conservative 0 /organism="unknown" /mol_type="unassigned DNA" from GI:12827623 GI:12827625 patent US 15 15 0; Score 15; DB Pred. No. 1.2 0; Mismatches 15 bp 6126965. 15 bp 6126965. DB 6; I 1.2e+03; DNA DNA Rahman, A. Length 15, Indels linear linear PAT 14-FEB-2001 PAT 14-FEB-2001 0 Gaps 0

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Semple, S.C., Klimuk, S.K., Harasym, T., Hope, Cullis, P., Scherrer, P. and Debeyer, D. Charged therapeutic agents encapsulated in containing four lipid components

Patent: US 6287591, A 15 11-SEP-2001;
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Sequence 1 from Patent WO2004017944.
CQ789712
CQ789712.1 GI:45823264
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                                                                                                                           Zhang, J.A. and Ahmad, I.
Liposomal gemcitabine compositions for
Patent: WO 2004017944-A 1 04-MAR-2004;
                                                                                                                                                                               synthetic construct synthetic construct
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                                                                                                                  Neopharm, Inc. (US)
                                                                                                                                                                    other sequences; artificial sequences.
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Location/Qualifiers
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15 from patent US 6287591.
                                                   /organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/note="Anti-raf-oligonucleotides"
                                                                                                       Location/Qualifiers
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/mol_type="unassigned
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/mol_type="unassigned/
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Kasid,U., Gokhale,P., Zhang,C., Dritschilo,A. and Rahman,A.
Cationic liposomal delivery system and therapeutic use thereof
Patent: US 6559129-A 1 06-MAY-2003;
Location/Qualifiers
                                                                                                                                                                                                                                                                 1 (bases 1 to 15)
Kasid, U., Gokhale, P., Zhang, C., Dritschilo, A. and Rahman, A.
Cationic liposomal delivery system and therapeutic use thereof
Patent: US 6559129-A 3 06-MAY-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AR310685
Sequence 1 from patent
AR310685
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                       25 from Patent W003039595
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/mol_type="genomic
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Pred. No. 1.2e+03;
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Inex Pharmaceuticals Corp. (CA)
Location/Qualifiers
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                                                                                                                                                            Sequence 25 from Patent
AX957740
AX957740.1 GI:40785558
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                              Homo sapiens (human)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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           Tam,Y.K., Semple,S., Klimuk,S. and Chikh,G. Cancer vaccines and methods of using the same Patent: WO 03094828-A 25 20-NOV-2003;
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Methylated immunostimulatory oligonucleotides
                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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Sequence
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                                                                              Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.

1 (bases 1 to 15)

Semple,S.C., Klimuk,S.K., Harasym,T., Hope,M.J., Ansel,S.M.,

Cullis,P., Scherrer,P. and Debeyer,D.S.

High efficiency encapsulation of charged therapeutic agents in
lipid vesicles
lipid vesicles
patent: JP 2002501511-A 15 15-JAN-2002;

INEX PHARMACEUTICALS CORP
PD 15-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15;
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STEVEN M ANSELL,
PI PIETER CULLI
CC Strandedness
                                                                                                                                                                                                                                                                                                                                             High efficiency encapsulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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 PIETER CULLIS, PETER SCHERRER, DAN SUITE DEBEYER Strandedness: Single;
                                               14-MAY-1998 JP 1998548646
SEAN C SEMPLE, SANDRA K KLIMUK, TROY HARASYM, MICHAEL J HOPE,
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
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Sequence 49 from patent US
AR216002
AR216002.1 GI:23314290
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Monia,B.P. and Boggs,R.T.
Antisense oligonucleotide modulation of Patent: US 5952229-A 14-SEP-1999;
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Monia, B.P.
Antisense oligonucleotide inhibition of raf gene expression Patent: US 6410518-A 49 25-JUN-2002;
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/mol_type="genomic DNA"
/db_xref="taxon:35827"
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/mol_type="unassigned DNA"
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Sequence 2 from
AR110776
                                            1 (bases 1 to 25)
Kasid,U., Gokhale,P., Zhang,C., Dritschilo,A. and Rahman,
Cationic liposomal delivery system and therapeutic use th
Patent: US 6559129-A 2 06-MAY-2003;
Location/Qualifiers
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1 (bases 1 to 25)

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Kasid, U., Gokhale, P., Dritschilo, A. an Liposomes containing oligonucleotides Patent: US 6126965-A 2 03-OCT-2000;

Location/Qualifiers
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/mol_type="genomic DNA"
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/mol_type="unassigned DNA"
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1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/PCT_NBW_PUB.seq:*
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Maximum Match 100%
Listing first 45 summaries
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Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
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gn2 6/ptodata/1/pubpna/USO8 NEW PUBCOMB.seq:*
gn2 6/ptodata/1/pubpna/USO8 PUBCOMB.seq:*
gn2 6/ptodata/1/pubpna/USO9 PUBCOMB.seq:*
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cgn2 6/ptodata/1/pubpna/USO9 PUBCOMB.seq:*
cgn2 6/ptodata/1/pubpna/USO9 NEW PUB.seq:*
cgn2 6/ptodata/1/pubpna/USO9 NEW PUB.seq:*
cgn2 6/ptodata/1/pubpna/USIOR PUBCOMB.seq:*
cgn2 6/ptodata/1/pubpna/USIOR PUBCOMB.seq:*
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6/ptodata/1/pubpna/US11A FUBCOMB.seq: *
6/ptodata/1/pubpna/US11 NEW PUB.seq: *
6/ptodata/1/pubpna/US60 NEW PUB.seq: *
[6/ptodata/1/pubpna/US60_PUBCOMB.seq: *
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 US-09-895-480A-15

US-10-290-545-25

US-10-365-623-16

US-10-075-994A-1

US-10-075-994A-3
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US-09-930-283A-3
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15.123 Million cell updates/sec
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Sequence 1, Appli
Sequence 3, Appli
Sequence 15, Appl
Sequence 25, Appl
Sequence 16, Appl
Sequence 1, Appli
Sequence 3, Appli
                                                                                                        Description
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US-09-930-283A-1
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	equence 6. Appl	quence 5, Appl	equence 6, Appl	e 5,	e 3, Appl	: 5, App	e 127444	127	12716	Ü	W	- 1	in	e 27212		_	o o	U	e 12	e 657, Ap	19,	equence 49	equence 2,	e 2, Appl	equence 817261	e 11354	e 113550	e 113549	equence 2,	quence 2, Ar	e 47, App	equence 49, App	equence 15, App	e 25, App	equence 25, App	equence 25,	equence 1, Appl	Sequence 4, Appli

ALIGNMENTS

ATTORNEY/AGENT INFORMATION:

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                                             PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/354,109
FILING DATE: 1999-07-15
ATTORNEY/AGENT INFORMATION:
NAME: Hendricke, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: kasid
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 591-4470
TELEPHONE: (703) 591-4428
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity luv.
15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 22031
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gokhale, Prafulla
Dritschilo, Anatoly
TITLE OF INVENTION: Liposomes containg Oligonucelotides
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: unknown
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
ANTI-SENSE: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Hendricks, Glenna
REGISTRATION NUMBER: 32,535
REFERENCE/DOCKET NUMBER: kasid
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 591-4470
            TOPOLOGY: unknown MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kasid, Usha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/930,283A FILING DATE: 16-Aug-2001 CLASSIFICATION: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: VA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: P.O. Box 2509
CITY: Fairfax
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (703)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DDRESSEE: Hendricks and Assoc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 15; DE 100.0%; Pred. No. 1.3 /ative 0; Mismatches
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RESULT 4 US-10-290-545-25

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; ANTI-SENSE: YES
; SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-930-283A-3
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US-09-895-480A-15
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                                                                 Matches
                                                                                Query Match
Best Local &
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APPLICANT: Inex Pharmaceuticals Inc.
TITLE OF INVENTION: High Efficiency Encapsulation of Charged Therapeutic
                                                                                                                                                                                                                                                                                       TELEFAX: <Unknown>
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/895,480A
FILING DATE: 29-Jun-2001
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
APPLICATION UNMBER: cUnknown>
FILING DATE: cUnknown>
ATTORNEY/ACENT INFORMATION:
NAME: cUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
HYPOTHETICAL: no
ANTI-SENSE: yes
                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: <Unknown>
REFERENCE, NUMBER: <Unknown>
TELECOMMUNICATION INFORMATION:
TELEPHONE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Word Perfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS 5.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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                     1 GTGCTCCATTGATGC 15
                                                                                 Similarity
GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 80435
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STATE: CO
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STREET: PO Box 5068
                                                                                                                                                                                                                                                                       LENGTH: 15
                                                                 Conservative
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                                                                                 Score 15; DB 10;
Pred. No. 1.3e+02;
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                                                                   Mismatches
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OTHER INFORMATION: Antisense oligonucleotide directed against human c-raf-1 protein TOTHER INFORMATION: kinase
US-10-365-623-16
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US-10-365-623-16
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                                                                                                                                                                                                                    Sequence 1, Application US/10075994A Publication No. US20030215489A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Klimuk, Sandy
APPLICANT: Yuan, Zuan-Ning
TITLE OF INVENTION: Improved Mucosal Vaccines and Methods for Using the
FILE REFERENCE: A-71854/TAL/AXG
CURRENT APPLICATION NUMBER: US/10/290,545
CURRENT FILING DATE: 2002-11-07
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin version 3.1
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Xie, Dong
TITLE OF INVENTION: Protein Carrier System for Therapeutic Oligonucleotides
FILE REFERENCE: 63024.000001
CURRENT APPLICATION NUMBER: US/10/365,623
CURRENT FILING DATE: 2003-02-13
CURRENT FILING DATE: 2003-02-13
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APPLICANT: Semple, Sean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25, Application US/10290545 Publication No. US20030125292A1
                                                                                                                                                                 APPLICANT: KASID, APPLICANT: GOKHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn version 3.1
SEQ ID NO 16
                                   FILE REFERENCE: 219604
                                        APPLICANT: RAHMAN, Aquilur
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
                                                                                                           APPLICANT:
                                                                                                                            APPLICANT: PIE,
APPLICANT: MEWAI
                 CURRENT APPLICATION NUMBER: US/10/075,994A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENGTH: 15
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                                                                                     MEWANI, Rajshree
AHMAD, Imran
DRITSCHILO, Anatoly
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                                                                                                                                                               GOKHALE, Prafulla
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                                                                                                                                                  Jin
                                                                                                                                                                                   Usha
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2002-02-15
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Pred. No.
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; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic
US-10-075-994A-3
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                                                                                                                                                                                                                                                                                                           US-10-075-994A-4
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FILE REFERENCE: 219604
CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.2
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.2 SEQ ID NO 3 LENGTH: 15
                                                                                                                                                                                                                                                      GENERAL INFORMATION
                                                                                                                                                                                                                                                                     Sequence 4, Application US/10075994A Publication No. US20030215489A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local Similarity
Matches 15; Conserv
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APPLICANT:
APPLICANT:
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                                                                                                      APPLICANT: MEWANI, Rajshree
APPLICANT: AHMAD, Imran
APPLICANT: DRITSCHIIO, Anatoly
APPLICANT: RAHMAN, Aquilur
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
                                                                                                                                                                                                                APPLICANT: KASID, APPLICANT: GOKHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 219604
CURRENT APPLICATION NUMBER: US/10/075,994A
CURRENT FILING DATE: 2002-02-15
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: DRITSCHILO, Anatoly
APPLICANT: RAHMAN, Aquilur
TITLE OF INVENTION: CHEMOSENSITIZING WITH LIPOSOMES CONTAINING OLIGONUCLEOTIDES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: KASID,
APPLICANT: GOKHAI
APPLICANT: PIE,
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SOFTWARE: PatentIn version 3.2
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ORGANISM: Artificial Sequence
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AHMAD, Imran
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RESULT 10
US-10-437-263-25
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US-10-347-924-1
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Publication No. US20040009943A1
GENERAL INFORMATION:
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TITLE OF INVENTION: PATHOGEN VACCINES AND METHODS FOR USING THE SAME FILE REFERENCE: A-72216/TAL CURRENT FILING DATE: 2003-05-12
PRIOR APPLICATION NUMBER: 60/379,343
PRIOR APPLICATION NUMBER: 60/379,343
PRIOR FILING DATE: 2002-05-10
PRIOR PRIOR PATHORATION NUMBER: 60/460,646
PRIOR PILING DATE: 2003-04-04
PRIOR APPLICATION NUMBER: 60/454,298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Gokhale, Prafulla
APPLICANT: Gokhale, Prafulla
APPLICANT: Zhang, Chuanbo
APPLICANT: Dristschilo, Anatoly
APPLICANT: Rahman, Aquilur
TITLE OF INVENTION: CATIONIC LIPOSOMAL DELIVERY SYSTEM AND THERAPEUTIC USE THEREOF
FILE REFERENCE: 220807
CURRENT APPLICATION NUMBER: US/10/347,924
CURRENT FILING DATE: 2003-01-21
PRIOR APPLICATION NUMBER: US 09/354,109
PRIOR FILING DATE: 1999-07-15
PRIOR APPLICATION NUMBER: US 08/957,327
PRIOR FILING DATE: 1997-03-21
PRIOR FILING DATE: 1997-03-21
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                                                                                                                                                                                                                         APPLICANT: Semple, Sean
APPLICANT: Tam, Ying K.
APPLICANT: Chikh, Ghania
APPLICANT: Hope, Michael J.
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Sequence 25, Application US/10437258

Publication No. US20040013649A1

GENERAL INFORMATION:
APPLICANT: Tam, Ying K.
APPLICANT: Tam, Ying K.
APPLICANT: Chikh, Ghania
APPLICANT: Chikh, Ghania
FILE OF INVENTION: CANCER VACCINES AND METHODS OF USING THE SAME
CURRENT APPLICATION NUMBER: US/10/437,258

CURRENT APPLICATION NUMBER: 60/379,343

PRIOR APPLICATION NUMBER: 60/460,646

PRIOR APPLICATION NUMBER: 60/460,646

PRIOR APPLICATION NUMBER: 60/460,646

PRIOR FILING DATE: 2003-04-04

PRIOR APPLICATION NUMBER: 60/454,298
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-437-263-25
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US-10-437-275-25
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PRIOR FILING DATE: 2002-05-10
PRIOR APPLICATION NUMBER: 60/460,646
PRIOR FILING DATE: 2003-04-04
PRIOR APPLICATION NUMBER: 10/290,545
PRIOR FILING DATE: 2002-11-07
NUMBER OF SEQ ID NOS: 32
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SEQ ID NO 25
LENGTH: 15
TYPE: DNA
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NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn version 3.2
SEQ ID NO 25
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APPLICANT: Semple, Sean
APPLICANT: Semple, Sean
APPLICANT: Klimuk, Sandra
APPLICANT: Chikh, Ghania
ITITLE OF INVENTION: USING THE SAME
FILE REFERENCE: A-72158/TAL
CURRENT APPLICATION NUMBER: US/10/437,275
CURRENT FILING DATE: 2003-05-12
CURRENT FILING DATE: 2003-05-12
CURRENT FILING DATE: 2003-05-12
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SEQ ID NO 25

PRIOR FILING DATE: 2003-03-12 NUMBER OF SEQ ID NOS: 34 SOFTWARE: PatentIn version 3.2

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RESULT 13
US-10-925-734-15
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                                                                                                       US-10-925-734-15
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                         Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15, Application US/10925734 Publication No. US20050008689A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                            TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Inex Pharmaceuticals Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 15
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 80435

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.4

MEDIUM TYPE: Diskette, 3.5 inch, 1.4

COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS 5.0

SOFTWARE: WOXT PERFECT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0/925,734

FILING DATE: 24-Aug-2004
CLASSIFICATION: «UDKNOWN»
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/895,480

FILING DATE: 29-JUN-2001
ATTORNEY/ACENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: <Unknown>
REFERENCE/DOCKET NUMBER: <Unknown>
TELECOMMUNICATION INFORMATION:
                                                                                                                  HYPOTHETICAL: no
ANTI-SENSE: yes
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: other nucleic
                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: High Efficiency Encapsulation of Charged
1 GTGCTCCATTGATGC 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                     TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                            TELEPHONE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Dillon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: PO Box 5068
                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
                                                                                                                                                                                                                                         LENGTH: 15
                                                                                                                                                                                                                                                                                                         TELEFAX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                   NAME: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Oppedahl & Larson LLP
                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 15; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Therapeutic
                                0
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                                               Score 15; DB 21;
Pred. No. 1.3e+02;
                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                    acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 15;
                                                               Length 15;
                                Indels
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                                0;
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                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               밁
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NUMBER OF SEQ ID NOS: 109
SEQ ID NO 47
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 47, Application US/10173225B Publication No. US20030119769A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local (
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SEQ ID NO 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: US 10/057,550
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: US 09/143,214
PRIOR FILING DATE: 1998-08-28
PRIOR APPLICATION NUMBER: PCT/US98/13961
PRIOR FILING DATE: 1908-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-06-07-07-06-07-06-07-06-07-06-07-07-07-06-07-07-07-07-07-07-07-07-07-07-
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PRIOR APPLICATION NUMBER: 09/506,073
PRIOR FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: US 09/143,214
                                                                                                                          PRIOR FILING DATE: 1995-05-31
PRIOR APPLICATION NUMBER: US 08/250,856
PRIOR FILING DATE: 1994-05-31
                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/US95/07111
                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 08/756,806 PRIOR FILING DATE: 1996-11-26
                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1998-07-06
PRIOR APPLICATION NUMBER: US 08/888,982
PRIOR FILING DATE: 1997-07-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Antisense Oligonucleotide Modulation
FILE REFERENCE: ISPH-0665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 08/756,806 PRIOR FILING DATE: 1996-11-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1998-07-06
PRIOR APPLICATION NUMBER: US 08/888,982
PRIOR FILING DATE: 1997-07-07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/250,856 FILING DATE: 1994-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/US95/07111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 1995-05-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/10/173,225B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GTGCTCCATTGATGC 15
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15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1998-08-28
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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; TYPE: DNA; OF TYPE: DNA; OF TYPE: DNA; OF TYPE: DNA; OF TYPE: OF TYPE: TYPE:
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                      Minimum DB seq length: 0 Maximum DB seq length: 50
                                                                                                                                                                                                                                                                          Searched:
                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                          Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    OM nucleic - nucleic search, using sw model
                                                                                                                          Database :
                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                             Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                             Run on:
IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                          US-10-075-994A-1
15
                                                                                                                                                                                                                                                                                                                                                                                                June 21, 2005, 05:02:47; Search time 2999 Seconds (without alignments)
190.385 Million cell updates/sec
                                                                                                                                                                                                                                                                           34239544 seqs, 19032134700 residues
                                                                                                                                                                                                                                                                                                                                            1 gtgctccattgatgc 15
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 gb est1:
gb htc: *
gb htc: *
gb est3: *
gb est4: *
gb est4: *
gb est6: *
gb est6: *
gb est6: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
ŀ	12.4	82.7	40	μį	AA434107	AA434107 zw24b01.s
ი 2	12.4	82.7	41	9	CL213712	A037F07
	12.4		41	9	CW509288	
	12.4	82.7	43	9	CC894144	CC894144 RRK138 Ba
	12.4	82.7	47	8	CC200383	RRE120
	11.8	78.7	32	9	CL659092	CL659092 PRI0133a
7	11	73.3	27	9	AG190569	AG190569 Pan trogl
8	11	73.3	39	8	AZ579181	AZ579181 1M0363G10
9	11	73.3	46	φ	CL982908	CL982908 GC0134 TI
c 10	11	73.3	49	9	BX659267	BX659267 Arabidops
11	10.8	72.0	26	8	AZ769902	AZ769902 1M0571G06
	10.8	72.0		Н	AU256788	AU256788 AU256788
c 13	10.8	72.0	28	в	AZ780454	AZ780454 2M0017016
	10.8	72.0	42	9	TA59H06P	AL455680 T. brucei
15	10.8	. 72.0	43	7	CV521218	CV521218 0089P0056
16	10.8	72.0	43	9	CL256795	CL256795 XT0290 Sa
c 17	10.8	72.0		œ	AZ317769	AZ317769 1M0036002
18	10.8	72.0		μ	AI941313	AI941313 sb87e01.y
	10.8	72.0	47	7	CF339432	CF339432 RCL104-
c 20	10.8	72.0	49	ب	AA444355	AA444355 vd92d04.r
21	10.8	72.0	50	μ	AJ558646	AJ558646 AJ558646
c 22	10.8	72.0	50	بر	AU103457	AU103457 AU103457
c 23	10.8	72.0	50	σ	CB226725	CB226725 1Ru29D04
24	10.4	69.3	22	9	TA197A04Q	AL477173 T. brucei

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10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.2	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4	10.4
68.0	68.0	68.0	68.0	68.0	68.0	68.0	68.0	68.0	68.0	68.0	69.3	69.3	69.3	69.3	69.3	69.3	69.3		69.3	69.3
43	י ה י ה	43	41	40	35	3 5	34	ω ω	27	26	49	49	49	49	44	38	34	31.	30	23
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BJ001161	AV833106	AI426076	AL946655	AI900047	CG785587	AZ654998	CC049952	ВJ060580	AZ991985	AZ308069	BX654661	BH852072	AI888465	AI035265	CG781920	BZ664429	AZ786136	AG199008	BZ596965	AZ760244
			AL946655 Arab	AI900047 sb98	CG785587 00S0	AZ654998 1M05	CC049952 01S0	BJ060580 BJ06	AZ991985 2M02	AZ308069 1M00	BX654661 Arab	BH852072 SALK	AI888465 wn32	AI035265 ue18	CG781920 1123	BZ664429 SALK	AZ786136 1M00	AG199008 Pan	BZ596965 SALK	AZ760244 1M05
)1161	901EEBAR	mh31h05.x	Arabidops	sb98b08.y	00S0119-0	1M0529L12	01S0530-0	ВJ060580	2M0276P21	1M0010N09	Arabidops	SALK_0741	wn32a07.x	ue18b08.x	1123047D1	SALK_0711	1M0031M21	Pan trogl	SALK_0979	1M0553C19

ALIGNMENTS

	FEATURES FEATURES BOUICE FEATURES	AUTHORS H M TITLE W JOURNAL U COMMENT C	3	RESULT 1 AA434107/c LOCUS DEFINITION Z I
/organism="Homo sapiens" /mol type="mRNA" /mol type="mRNA" /db_xref="mRNA" /db_xref="mRNA" /db_xref="mRNA" /db_xref="taxon:9606" /db_xref="taxon:9606" /clone="InAGE:770185" /tissue_type="ovarian tumor" /tissue_type="ovarian tumor" /tissue_type="ovarian tumor" /lab host="DH10B (ampicillin registant)" /clone_lib="Soares ovary tumor NDHOT" /note="Organ: ovary; Vector: pT773D (Pharmacia) with a modified polylinker; Site 1: Not I; Site 2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5'	washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 Tel: 314 286 1800 Fax: 314 286 1810 Email: est@watson.wustl.edu This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. Trace considered overall poor quality Possible reversed clone: similarity on wrong strand Seq primer: -41m13 fwd. ET from Amersham High quality sequence stop: 1. Location/Qualifiers 140	Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R. and Wilson, R. WashU-Merck EST Project 1997 Unpublished (1997) Contact: Wilson RK	GI:2139021 s (human) s (human) Metazoa; Chordata; Craniata; Vert utheria; Primates; Catarrhini; Ho to 40)	AA434107 40 bp mRNA linear EST 29-MAY-1997 EW24b01.s1 Soares ovary tumor NbHOT Homo sapiens cDNA clone IMAGE:770185 3' similar to TR:G556217 G556217 SPLICEOSOMAL PROTEIN.

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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     analysis of the mouse genome Proc. Natl. Acad. Sci. U.S.A. 100 (17), 9918-9922 (2003)
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Hansen,J., Floss,T., van Sloun,P., Fuchtbauer,E.M., Vauti,F.,
Arnold,H.H., Schnutgen,F., Wurst,W., Von Melchner,H. and Ruiz,P.
Alarge-scale, gene-driven mutagenesis approach for the functional
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           German Genetrap Consortium (GGTC) Email: info@genetrap.de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         On Jun 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                          /cell_type="mmbryonic stem cell"
/cell_line="ES cells 12952 (former)
/clone_lib="GGTC Gene Trap Library
/note="Vector: pTlbetageo"
                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="A037F07"
                                                                                                                                                                                                                                                                                                                                                                                                                                           mol_type="mRNA"
strain="129 Sv"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          organism="Mus musculus"
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                                                                                                                                                                                                                                                                                                                                                                               sex="Male"
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92.9%;
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Pred. No. 2e+04;
0; Mismatches 1;
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Pred. No. 2e+04;
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                                                                   On Sep 2, 2003 this sequence version replaced gi:33392557. Contact: BayGenomics
Contact: BayGenomics
Bay Area Functional Genomics Consortium (BayGenomics)
Bmail: info@baygenomics.ucsf.edu
Sequence tag generated by 5' RACE of total RNA from gene trap Ex
Sequence tag generated by 5' RACE of total RNA from gene trap Ex
cell line. ES cell lines harboring insertion mutation of target
gene are available upon request from BayGenomics. Annotation
information available from
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Sequence tag generated by 5' RACE of total RNA from gene trap E
cell line. ES cell lines harboring insertion mutation of target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http://baygenomics.ucsf.edu/
Unpublished (2001)
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CW509288
                Class: Gene Trap
                                  CELL_LINE&KEY=RRK138
                                                       http://baygenomics.ucsf.edu/cgi-bin/BaySearch.py?OPTION=EXACT&TYPE=
                                                                                                                                                                                                                                 http://baygenomics.ucsf.edu/
Unpublished (2001)
                                                                                                                                                                                                                                                                          BayGenomics.
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 43)
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CC894144.2
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RRK138 BayGenomics Gene Trap
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Bay Area Functional Genomics Consortium (BayGenomics)
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/note="Vector: pGTlLxf"
Location/Qualifiers
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92.9%;
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RESULT 6
CL659092/c
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AUTHORS
TITLE
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CC200383/c
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Best Local (
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: BayGenomics
Contact: BayGenomics
Consortium (BayGenomics)
Bay Area Functional Genomics Consortium (BayGenomics)
Bay 1: info@baygenomics.ucsf.edu
Bequence tag generated by 5' RACE of total RNA from gene trap E8
Sequence tag generated by 5' RACE of total RNA from gene trap E8
cell line. ES cell lines harboring insertion mutation of target
gene are available upon request from BayGenomics. Annotation
information available from
CL659092 32 bp DNA linear GSS 09-JUL-20 PRI0133a_G11 - PRI0133a_B21.1 (32) Mixed stage fosmid library of pacificus var. California Pristionchus pacificus genomic, genomic survey sequence.
                                                                                                                                                      N
                                                                                                                                                                                       13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC200383 47 bp mRNA linear GSS 0:
RRE120 BayGenomics Gene Trap Library pGT2Lxf Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N
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Unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          http://baygenomics.ucsf.edu/cgi-bin/BaySearch.py?OPTION=EXACT&TYPE=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BayGenomics.
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strain="129 ola"
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                                                                                                                                                                                                                                                                                                                                                strain="129 ola"
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92.9%;
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92.9%;
                                                                                                                                                                                                                                                              _type="Embryonic stem cell"
= lib="BayGenomics Gene Trap Library pGT2Lxf"
="Vector: pGT2Lxf"
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Pred. No. 2e+04;
0; Mismatches 1;
                                                                                                                                                                                                  Score 12.4; DB 8;
Pred. No. 2.1e+04;
                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                   DB 8;
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                                              GSS 09-JUL-2004
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AG190569
LOCUS
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AUTHORS
TITLE
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                                                                                                                                       TITLE
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                                                                                                                                                                        AUTHORS
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CL659092.1
GSS.
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                                                                                                                                                                                                                   BAC end sequences of Library RP-43
                                                                                                                                                                                                                                                                                                               Pan troglodytes (chimpanzee)
Pan troglodytes
                                                                                                                                                                                                                                                                                                                                                            sequence.
AG190569
AG190569.1 GI:45222745
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                                                                                                                                                                                                  Unpublished
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Pristionchus
/mol_type="genomic DNA"
/strain="California"
/db_xref="taxon:54126"
                                                                                                                                                                                       1 to 27)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     clone lib="Mixed stage
ar. California"
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86.7%;
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Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea (B-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance of clone tracking errors.
                                                                                                                                                                                                                                                                     Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H. Direct Submission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Evolutionary Biology
Max-Planck-Institute for Developmental Biology
Spemannstr. 37-39, Tuebingen D-72076, Germany
Tel: 00497071601371
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Nucleic Acids Res. 32 (1), D421-D422 (2004)
Contact: Sommer RJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Park,H., Kim,Y., Kim,S., Han,Y., Woo,T., Park,K., Eun,C.J., Hoon,S.T., Chu,M., Kim,H., Joo,S., Kim,C., Song,W. and Yoo,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pan troglodytes DNA, clone: RP43-066B21.TJ, genomic survey
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This library was generated at Caltech, Pasadena, USA and end
sequenced at Vancouver, Canada.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Srinivasan, J., Otto, G.W., Kahlow, U., Geisler, R. and Sommer, R. AppaDB: an AcedB database for the nematode satellite organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pristionchus pacificus
Pristionchus pacificus
Eukaryota, Metazoa, Nematoda, Chromadorea, Diplogasterida,
Neodiplogasteridae, Pristionchus.
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Pred. No. 4.4e+04;
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RESULT 8
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                                                                                                                                                                                                                                                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0363 row: G column: 10
Seg primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wilght, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GTGCTCCATTG 11
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Unpublished (2000)
                                                                                                                                                                                                                                                                                           High quality sequence stop: 39.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                         Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus
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LIBRARY
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R.Site 1 : EcoRI
R.Site 2 : EcoRI
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                                                              /lab_host="B. Coli strain XL10-Gold, T1-resistant, F-"/clone_lib="Mouse_10kb_plasmid_UUGCIM_library"/note="Vector: pW042nv; Purified_genomic_DNA_from_M.musculus_C57BL/6J (male) was obtained_from_the_Jackson_musculus_C57BL/6J (male) was obtained_from_the_Jackson_musculus_C57BL/6J (male)
                                                                                                                                                                                                                                   /mol
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1. .27
                     Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/).
                                                                                                                                                  /sex≃"Male"
                                                                                                                                                                                        db_xref="taxon:10090"
                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                                                                    organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="male"
/cell_type="lymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mol_type="genomic DNA'
/db_xref="taxon:9598"
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                                                                                                                                                                     clone="UUGC1M0363G10"
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100.0%; Pr
hydrodynamically sheared by repeated passage through
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 46) Cobellis,G., Nicolaus,G., Marra,E., Barbarisi,M., Sardiello,M., Di Giorgio,F.P., Iovino,N., Zollo,M., Ballabio,A. and Cortese,R. Tagging genes with cassette-exchange sites Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                          Sequence tag generated by 5' RACE of total RNA from gene trap Education line. ES cell lines harboring insertion mutation of target gene are available upon request from TIGEM. Annotation informationable from TIGEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 TCCATTGATGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cobellis@tigem.it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fax: +390815790919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Via P. Castellino, 111, 80131 NAPOLI, ITALY
Tel: +390816132205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: TIGEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CL982908.1 GI:52420303
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                               Similarity
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                                                                                                                                                                                                                       /strain="129 ola"
/db_xref="taxon:10090"
/clone="8644.24"
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                                                                                                             /cell_type="Embryonic stem cell"
/cell_line="El4"
/clone lib="TIGEM gene trap library"
/note="Vector: pFLIP1"
                                                                                                                                                                                                                                                                                     organism="Mus musculus"
/mol_type="mRNA"
                                                                                                                                                                                                     /sex="male"
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                             73.3%;
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        <u>,</u>
                             Score 11;
Pred. No.
           Mismatches
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DB 9; LC
3. 1.4e+05;
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BX659267/c
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                                                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                Zuechtungsforschung, Carl-von-Linne-Weg 10, Koeln, 50829, Germany This sequence has been recovered from the left border of the T-DNA. It indicates an insertion close to or within gene Atl972180. Details on the protocols used for generation of the sequence are described in References 1-3. The sequences are generated at the MPI for Plant Breeding Research in the context of the GABI-Kat project. GABI-Kat is part of the German Plant Genomics program designated 'GABI'. Information on line availability can be found at: http://www.mpiz-koeln.mpg.de/GABI-Kat/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   High-throughput generation of sequence indexes from T-DNA mutagenized Arabidopsis thaliana lines BioTechniques 35 (6), 1164-1168 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             An Arabidopsis thaliana T-DNA mutagenized population (GABI-Kat) for flanking sequence tag-based reverse genetics Plant Mol. Biol. 53 (1-2), 247-259 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Li,Y., Rosso,M.G., Strizhov,N., Viehoever,P. and Weisshaar,B. GABI-Kat SimpleSearch: a flanking sequence tag (FST) database for the identification of T-DNA insertion mutants in Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (31-MAR-2004) Weisshaar B., Max-Planck-Institut fuer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Strizhov, N., Rosso, M.G., Li, Y. and Weisshaar, B.
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                   /note="PCR was performed on DNA from Arabidopsis thaliana plants (T1) which were transformed with the T-DNA from vector pAC161 (GenBank accession number: AJ537514). The lines contain one or more T-DNA insertions. The DNA fragment(s) resulting from the PCR were directly sequenced to determine the genomic sequence flanking the insertion.
                                                                                                                                                                                 ecotype="Col-0"
                                                                                                                                                                                                                                                                                                /mol_type="genomic D
/strain="Columbia 0"
                                                                                                                                                                                                                                                                                                                                                organism="Arabidopsis thaliana"
                                                                                                                                                                                                                clone_lib≃"Arabidopsis thaliana T-DNA insertion lines"
                                                                                                                                                                                                                                             clone="GK-644G02-022818"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Li,Y., Strizhov,N., Reiss,B., Dekker,K. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Li,Y., Rosso,M.G., Viehoever,P., Dekker,K.A.
                                                                                                                                                                                                                                                                _xref="taxon:3702"
derived sequences were removed
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Query Match Best Local Similarity

73.3%;

Score 11; DB 9; I Pred. No. 1.4e+05;

Length 49

Query Match Best Local Similarity

Score 10.8; DB 8; Pred. No. 1.7e+05;

Length 26;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D., Weiss,R.
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High quality sequence stop: 26.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Plate: 0571 row: G column: 06
Seq primer: CGTTGTAAAACGACGGCCAGT
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1M0571G06F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0571G06 F, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0571 row: G column: 06
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical Polymers Rese
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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            was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was but end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt end-sin high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                              /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/clone lib="Mouse 10kb plasmid UUGC1M library"/note="Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
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'clone="UUGC1M0571G06"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           strain="C57BL/6J"
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                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (Dases 1 to 28)
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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28 bp DNA linear GSS 16 2M0017016R Mouse 10kb plasmid UUGC1M library Mus musculus clone UUGC2M0017016 R, genomic survey sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Kikuya Kato
Graduate School of Biological Sciences
Nara Institute of Science and Technology
8916-5 Takayama, Ikoma, Nara 630-0101, Japan
Tel: 81-743-72-5581
University of Utah Genome Center University of Utah
                                                plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                 Mus musculus (house mouse)
                                                                                                                                                                                                                                                  AZ780454.1 GI:12912132 GSS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 28) Kato, K. and Matoba, R.
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                                Contact: Robert B. Weiss
                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
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/clone_Tib="3'-directed mouse cDNA library"
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/db_xref="taxon:10090"
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Insert Length: 10000 Std Error: 0.00
Plate: 0017 row: O column: 16
Seq primer: CACACAGAAACAGCTATGACC
Class: plasmid ends
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Fax: 801 585 7177
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Location/Qualifiers
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                                                       Conservative
                                                                                                                                                                                 was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pND42 (gi 4732114 | gb | AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource
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/strain="C57BL/6J"
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                                                                                 72.0%;
85.7%;
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                                                                                 Score 10.8; DB 8; Pred. No. 1.7e+05;
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23 GTGCTCCAGTTATG 10

TA59H06P

brucei sheared genomic

DNA ģ clone

59h06,

forward

sequence,

GSS 13-DEC-2000

linear

Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and nhl@sanger.ac.uk Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G. Direct Submission Trypanosoma brucei Trypanosoma brucei AL455680.1 GI:11857958 GSS. genomic survey sequence AL455680 Constructed at the Institute for Genomic Research Rockville, MD. Genomic DNA isolated from a cloned Trypanosoma. 1 (bases 1 to 42) Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae;

(TIGR), population

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Best Local S
  Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                source
                                                                                                                                                                                                                                                                                                                       Contact: Willis J
Department of Biology
Duke University
072-A Biological Sciences Science Drive, Durham, NC 27708, USA
Tel: 919 660 7340
Fax: 919 660 7293
Email: 941011s@duke.edu
Plate: 0089P0056 row: 02 column: D
Seq primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 GTGCCCCATTGAAG 9
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4 kb). The v + i method used for the library construction is
described in detail in Smith, H. and Venter, J.C. (Making small
insert libraries for whole genome shotgun sequencing projects. In
Genome Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barrell, Oxford University Press, 1999).
Email: nelsayedetigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        43 bp mRNA linear EST 06-OCT-2004 0089P0056Z.x0_D02 Mimulus guttatus library 2 Mimulus guttatus cDNA clone 0089P0056Z.x0_D02, mRNA sequence.
  12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; saterids; lamiids; Lamiales; Lamiales incertae sedis; Mimuleae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mimulus guttatus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Willis, J., Vision, T., Mimulus guttatus cDNA Unpublished (2004)
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Similarity 85.7%;
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|mol type="genomic DNA"
|strāin="TREU927"
|db xref="taxon:5691"
|clone="59h06"
                                                                                                                    /organism="Mimulus guttatus"
|mol_type="mRNA"
|mol_type="mRNA"
|db_xref="taxon:4155"
|/db_xref="taxon:4155"
|/clone="0089P0056Z.x0_D02"
|/clone_lib="Mimulus guttatus library 2"
|/note="Vector: pGEM-T Easy; a Mimulus guttatus cDNA
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Pred. No. 1.8e+05;
0; Mismatches 2;
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Pred. No. 1.8e+05;
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30 GIGTICGATIGATG 43
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Search completed: June 21, 2005, 06:51:11 Job time : 3008 secs

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Gapop 10.0 , Gapext 1.0
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184.004 Million cell updates/sec
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Copyright (c) 1993 - 2005 Compugen Ltd.
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CG531606 OST1114549
CG497924 OST38825
AA306005 EST177032
CG605299 OST281834
CG5601765 OST2413394
CG5601765 OST4413394
CG5604359 OST451337
CG54646546 OST144106
BF468476 UII-M-CD0-
AA332421 EST36381
CG577036 OST212398
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CG513061 OST66238
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ALIGNMENTS

JOURNAL COMMENT SOURCE ORGANISM RESULT 1 CG531606/c LOCUS ORIGIN REFERENCE FEATURES KEYWORDS VERSION ACCESSION DEFINITION Matches Query Match Best Local Similarity TITLE AUTHORS source Mammalia; Eutheria; Rodentia; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. E 1 (bases 1 to 165) Zambrowicz B.P., Abuin, A., Ramirez-Solis, R., Richter, L.J., Piggott, J., BeltrandelRio, H., Buxton, E.C., Edwards, J., Finch, R.A., Priddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B.W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markesich, D., Payne, R., Potter, D.G., Qian, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Zhu, Q., Person, C. and Sands, A.T. What kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention L. Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003) 15; CG531606 165 bp mRNA linear GSS 01-OCT-2003 OST114549 Mus musculus 129Sv/Ev Mus musculus cDNA clone OST114549, mRNA sequence. CG531606.1 GI:37318178 GSS. Email: materials@lexgen.com Gene trap sequence tag generated by 3' Class: Gene Trap. Contact: Zambrowicz BP Mus musculus described in Zambrowicz et al Lexicon Genetics Incorporated musculus (house mouse) 100.0%; Score 15; DB 9; llarity 100.0%; Pred. No. 7.3e+02; Conservative 0; Mismatches 0; /organism="Mus musculus" /mol_type="mRNA" /strain="129Sv/Ev" /db_xref="taxon:10000" /clone="0ST114549" /cell_type="embryonic stem cell" /clone_lib="Mus musculus 129Sv/Ev" location/Qualifiers . 165 (Nature. RACE from mouse ES cells as 1998 Apr 9;392(6676):608-11) Length 165; Indels TX 77381, USA °. Gaps 0

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1 GTGCTCCATTGATGC 15

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GIGCICCATIGATEC 25

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RESULT 3
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 207)

Adams, M.D., Kerlavage, A.R., Fleischmann, R.D., Fuldner, R.A., Bult, C.J., Lee, N.H., Kirkness, E.F., Weinstock, K.G., Gocayne, J.D., White, O., Sutton, G., Blake, J.A., Brandon, R.C., Man-Wai, C., Clayton, R.A., Cline, T.R., Cotton, M.D., Barle-Hughes, J., Fine, L.D., Fitzgerald, L.M., Fitzhugh, W.M., Fritchman, J.L., Geoghagen, N.S.,
                                                                                                                                                                                                                                                                           AA306005 207 bp mRNA linear E
EST177032 Jurkat T-cells VI Homo sapiens cDNA 5' end
proto-oncogene raf, mRNA sequence.
AA306005
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OST38825 Mus musculus 1295v/Ev N
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CG497924.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene trap sequence tag generated by 3' RACE from mouse ES cells as described in Zambrowicz et al (Nature. 1998 Apr 9;392(6676):608-11)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lexicon Genetics Incorporated
4000 Research Forest Drive, The Woodlands, TX 77381, USA
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                                                                                                                                                                                                                               Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: materials@lexgen.com
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/mol_type="mRNA"
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/clone="OST38825"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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similar to
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VERSION
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CG605299/c
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. E. (bases I to 229)

E. 1 (bases I to 229)

S. Zambrowicz, B.P., Abuin, A., Ramirez-Solis, R., Richter, L.J., Piggott, J., BeltrandelRio, H., Buxton, E.C., Edwards, J., Finch, R.A., Friddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Friddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B.W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markesich, D., Payne, R., Potter, D.G., Qian, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Shu, Q., Person, C. and Sands, A.T.

Zhu, Q., Person, C. and Sands, A.T.

Whkl kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
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OST281834 Mus musculus 129Sv/Ev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: arkerlav@tigr.org
For clone availability, additional sequence and expression
information related to this EST, please check the TIGR Human Gene
Index (http://www.tigr.org/tdb/hgi/hgi.html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other ESTs: THC192945
Contact: Kerlavage, AR
                                                                                                                                                                                                                                                                                                                                                                                                                                          GSS.
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                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CG605299.1 GI:37427977
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 3018699056
Fax: 3018699423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The Institute for Genomic Research 9712 Medical Center Drive, Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bioinformatics
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/db_xref="ATCC (inhost):160136"
/db_xref="taxon:9606"
/cell type="T-lymphocyte"
/clone_lib="Jurkat T-cells VI"
/note="Vector: pBluescript SK-; Site_1: EcoRI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/mol_type="mRNA"
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Pred. No. 7.
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s 0;
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CG661765/c
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Best Local S
Matches 15
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B. I (bases 1 to 23)

S. Zambrowicz, B.P., Abuin, A., Ramirez-Solis, R., Richter, L.J., Piggott, J., BeltrandelRio, H., Buxton, E.C., Edwards, J., Finch, R.A., Priddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B.W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markesich, D., Payne, R., Potter, D.G., Qian, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Zhu, Q., Person, C. and Sands, A.T.

Whkl kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention the proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
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     l Similarity 100
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CG661765.1
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OST443394 Mus musculus 1298v/Ev Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lexicon Genetics Incorporated 4000 Research Forest Drive, The Woodlands, TX 77381, USA Email: materials@lexgen.com
                                                                                                                                                                                                                                                                                      Email: materials@lexgen.com
Gene trap sequence tag generated by 3'
described in Zambrowicz et al (Nature.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GTGCTCCATTGATGC 15
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                                                                                                                                                                                                                                                                     Class: Gene Trap.
                                                                                                                                                                                                                                                                                                                                              Lexicon Genetics Incorporated
4000 Research Forest Drive, The Woodlands, TX 77381, USA
                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Zambrowicz BP
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                                                                                                                                      /db_xref="taxon:10090"
/clone="OST443394"
                                                                                                                                                                          /mol_type="mRNA"
/strain="129Sv/Ev"
                                                                                                                                                                                                                                                   ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /cell_type="embryonic stem cell"
/clone_lib="Mus musculus 129Sv/Ev"
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|mol_type="mRNA"
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100.0%; Sc.
100.0%; Pr
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                                                                                                _type="embryonic stem cell"
e_lib="Mus musculus 129Sv/Ev"
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   Score 15; DB 9; I
Pred. No. 7:7e+02;
; Mismatches 0;
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Pred. No. 7.7e+02;
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1998
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     Indels
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CG580338/c
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   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 274)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V.,
                                                                                                                                                                            BF897689.1
EST.
                                                                                                                                                                                                                                                                                                                                                                        1 GTGCTCCATTGATGC 15
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                                                                                                                                        Homo sapiens
                                                                                                                                                        Homo sapiens (human)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Class: Gene Trap.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lexicon Genetics Incorporated
4000 Research Forest Drive, The Woodlands,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CG58033
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OST219649 Mus musculus 129Sv/Ev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Email: materials@lexgen.com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Zambrowicz BP
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/strain="129Sv/Ev"
/db_xref="taxon:10090"
/clone="OST219649"
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100.0%; Pred. No. 7.9e+02;
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BF897689 274 bp mRNA linear QV1-MT0224-281100-512-b01 MT0224 Homo sapiens cDNA, BF897689
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Zambrowicz,B.P., Abuin,A., Ramirez-Solis,R., Richter,L.J., Piggott,J., BeltrandelRio,H., Buxton,E.C., Edwards,J., Finch,R.A., Priggott,J., Gupta,A., Hansen,G., Hu,Y., Huang,W., Jaing,C., Key,B., Jr., Kipp,P., Kohlhauff,B., Ma,Z.-Q., Markesich,D., Payne,R., Potter,D.G., Qian,N., Shaw,J., Schrick,J., Shi,Z.-Z., Sparks,M.J., Van Sligtenhorst,I., Vogel,P., Walke,W., Xu,N., Zhu,Q., Person,C. and Sands,A.T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene trap sequence tag generated by 3' RACE from mouse ES cells as described in Zambrowicz et al (Nature. 1998 Apr 9;392(6676):608-11)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wnk1 kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
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/clone_lib="Mus musculus 129Sv/Ev"
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                                                                             Piggott, J., BeltrandelRio, H., Buxton, E.C., Edwards, J., Finch, R.A., Friddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B.W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markesich, D., Payne, R., Potter, D.G., Qian, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Shu, Q., Person, C. and Sands, A.T. Whi, Q., Person, C. and Sands, A.T. Whi, Rinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003) Contact: Zambrowicz BP
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Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 (Dases 1 to 2/6)
Zambrowicz,B.P., Abuin,A., Ramirez-Solis,R., Richter,L.J.,
Direction of the control of the c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 276)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA sequence.
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281100-512-b01&t3=2000-11-28&t4=1)
Seq primer: puc 18 forward
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
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Lexicon Genetics Incorporated
                                                       OmniBank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: +55-11-2704922
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Location/Qualifiers
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/dev_stage="Adult"
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tive 0; Mismatch
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1 GTGCTCCATTGATGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zambrowicz,B.P., Abuin,A., Ramirez-Solis,R., Richter,L.J., Piggott,J., BeltrandelRio,H., Buxton,B.C., Edwards,J., Finch,R.A., Friddle,C.J., Gupta,A., Hansen,G., Hu,Y., Huang,W., Jaing,C., Key,B.W. Jr., Kipp,P., Kohlhauff,B., Ma,Z.-Q., Markesich,D., Payne,R., Potter,D.G., Qian,N., Shaw,J., Schrick,J., Shi,Z.-Z., Sparks,M.J., Van Sligtenhorst,I., Vogel,P., Walke,W., Xu,N., Sparks,M.J., Van Sligtenhorst,I., Vogel,P., Walke,W., Xu,N., Shu,Q., Person,C. and Sands,A.T.
Wnkl kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CG546546
CG546546.1 GI:37333133
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OST146106 Mus musculus 129Sv/Ev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: materials@lexgen.com Gene trap sequence tag generated by 3' RACE from mouse ES cells as described in Zambrowicz et al (Nature. 1998 Apr 9;392(6676):608-11) Class: Gene Trap.
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                                                                                                                                                                                                                                                                                                                                                     Class: Gene Trap.
                                                                                                                                                                                                                                                                                                                                                                        Gene trap sequence tag generated by 3' RACE from mouse ES cells as described in Zambrowicz et al (Nature. 1998 Apr 9;392(6676):608-11)
                                                                                                                                                                                                                                                                                                                                                                                                                 Email: materials@lexgen.com
                                                                                                                                                                                                                                                                                                                                                                                                                                  Lexicon Genetics Incorporated
4000 Research Forest Drive, The Woodlands, TX 77381,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Zambrowicz BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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                                                                                                                                                                                                                           /mol_type="mRNA"
/strain="1295v/Ev"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                                                   organism="Mus musculus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /cell_type="embryonic stem cell"
/clone_lib="Mus musculus 129Sv/Ev"
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/mol_type="mRNA"
/strain="129Sv/Ev"
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                                                                                                                                                                /cell_type="embryonic stem cell"
/clone_lib="Mus musculus 129Sv/Ev"
                                                                                                                                                                                                        /clone="OST146106"
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                                                                                   100.0%; Score 15; DB 9; 100.0%; Pred. No. 8e+02;
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Pred. No. 8e+02;
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AA332421/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA332421 294 bp mRNA 1
EST36381 Embryo, 8 week I Homo sapiens cDNA
proto-oncogene raf, mRNA sequence.
AA332421
                                                                                                                                                                   1 GTGCTCCATTGATGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDNA Library Preparation: M.B. Soares Lab Clone distribution: Researchers may obtain BMAP cDNA clones from RESEARCH GENETICS. It should be noted that Bento Soares is generating a small number of additional specialized non-redundant arrays of BMAP cDNAs whose availability will be considered under appropriate and limited collaborative arrangements The following repetitive elements were found in this cDNA sequence: 60-91, >GC_rich#Low_complexity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genome Res. 6 (9), 791-806 (1996) 97044477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP468476

293 bp mRNA linear EST 04-DEC-
UI-M-CD0-ayj-e-12-0-UI.rl NIH BMAP_Ret2 Mus musculus cDNA clone
UI-M-CD0-ayj-e-12-0-UI 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tel: 301 443 1706
Fax: 301 443 9890
Email: mEST@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bonaldo, M.F., Lennon, G. and Soares, M.B. Normalization and subtraction: two appr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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A 5' end
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RESULT 12
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Initial assessment of human gene diversity and expression patterns based upon 83 million nucleotides of cDNA sequence
AL Nature 377 (6547 Suppl), 3-174 (1995)
                                                                                                         mRNA sequence.
CG577036
CG577036.1 GI
GSS.
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Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                             CG577036 GSS 02-OCT-2003 OST212398 Mus musculus 129Sv/Ev Mus musculus cDNA clone OST212398,
                                                                                                                                                                                                                                                                                                                                                                                                                              1 GIGCICCATIGATGC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              For clone availability, additional sequence and expression information related to this EST, please check the TIGR Human Index (http://www.tigr.org/tdb/hgi/hgi/html)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The Institute for Genomic Research 9712 Medical Center Drive, Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bioinformatics
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Contact: Kerlavage, AR
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1 (bases 1 to 294)
                                                                                Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fax: 3018699423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer: M13 Reverse
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/clone_lib="Embryo, 8 week I"
/note="Organ: Embryo, 8 weeks; Vector: pBluescript SK-;
Site_1: EcoRI; Site_2: XhoI"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organisma"Homo sapiens"
/mol_type="mRNA"
/db_xref="ATCC (inhost):134082"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ocation/Qualifiers
                                                                                                                                        GI:37367640
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Pred. No. 8.1e+02;
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CG591346/c
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                                                                                                                                                                                                                                                                   Zambrowicz, B.P., Abuin, A., Ramirez-Solis, R., Richter, L.J., Piggott, J., BeltrandelRio, H., Buxton, E.C., Edwards, J., Finch, R.A., Friddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B.W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markesich, D., Payne, R., Potter, D.G., Qian, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Shuks, M.J., Person, C., and Sands, A.T.
Wnkl kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention Proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
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4000 Research Forest Drive, The Woodlands,
                                                                          Class: Gene Trap.
                                                                                                  Gene trap sequence tag generated by 3' RACE from mouse ES cells as described in Zambrowicz et al (Nature. 1998 Apr 9;392(6676):608-11)
                                                                                                                                                 Email: materials@lexgen.com
                                                                                                                                                                         Lexicon Genetics Incorporated
4000 Research Forest Drive, The Woodlands,
                                                                                                                                                                                                                                                      Contact: Zambrowicz BP
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/mol_type="mRNA"
/organism="Mus musculus"
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/db_xref="taxon:10090"
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Zambrowicz, B.P., Abuin, A., Ramirez-Solis, R., Richter, L.J., Piggott, J., BeltrandelRio, H., Buxton, B.C., Edwards, J., Finch, R.A., Friddle, C.J., Gupta, A., Hansen, G., Hu, Y., Huang, W., Jaing, C., Key, B. W. Jr., Kipp, P., Kohlhauff, B., Ma, Z.-Q., Markestch, D., Payne, R., Potter, D.G., Olan, N., Shaw, J., Schrick, J., Shi, Z.-Z., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Sparks, M.J., Van Sligtenhorst, I., Vogel, P., Walke, W., Xu, N., Shu, Q., Person, C. and Sands, A.T.

Zhu, Q., Person, C. and Sands, A.T.

Whil kinase deficiency lowers blood pressure in mice: a gene-trap screen to identify potential targets for therapeutic intervention proc. Natl. Acad. Sci. U.S.A. 100 (24), 14109-14114 (2003)
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CN295726
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CG513061.1 GI:37299634
GSS.
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                                      CN295726
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: materials@lexgen.com
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4000 Research Forest Drive, T
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/mol_type="mRNA"
/strain="129Sv/Ev"
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/strain="129Sv/Ev"
/db_xref="taxon:10090"
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/clone_lib="Mus musculus 129Sv/E
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/clone_lib="Mus musculus 129Sv/Ev"
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/clone="OST66238"
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Pred. No. 8.1e+02;
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                  cDNA 5', mRNA sequence.
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Search completed: June 21, 2005, 03:49:24
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278 GTGCTCCATTGATGC 264
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230 Constitution Drive, Menlo Park, CA 94025, USA
Tel: 650 473 8658
Fax: 650 473 7760
Email: rbrandenberger@geron.com
Insert Length: 309 Std Error: 0.00.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Brandenberger,R., Wei,H., Zhang,S., Lei,S., Murage,J., Figk,G.J., Li,Y., Xu,C., Fang,R., Guegler,K., Rao,M.S., Mandalam,R., Lebkowski,J and Stanton,L.W.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 309)
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EST.
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/note="oligo dT primed, full-length enriched cDNA library
from undifferentiated hES cell lines H1 (p32), H7 (p29),
and H9 (p26) maintained in feeder-free conditions"
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/mol_type="mRNA"
/db_xref="taxon:9606"
/tissue_type="embryonic stem cells, cell lines H1, H7, and
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Perfect score:
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Listing first 45 summaries
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Copyright (c) 1993 - 2005 Compugen Ltd.
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AR116777
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CQ789712
AR310685
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ALIGNMENTS

RESULT 2 ARI10777/c LOCUS LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE	Best Loca Matches Qy Db	SOURCE U ORGANISM U REFERENCE 1 AUTHORS K TITLE L JOURNAL P FEATURES SOURCE ORIGIN Query Match	RESULT 1 AR110775 LOCUS DEFINITION ACCESSION VERSION KEYWORDS
ARI10777 Sequence 3 from patent US 6126965. ARI10777 ARI10777.1 GI:12827625 Unknown. Unknown. Unclassified. 1 (bases 1 to 15) Kasid, U., Gokhale, P., Dritschilo, A. and Rahman, A. Liposomes containing oligonucleotides	Similarity 100.0%; Pred. No. 1.2e+03; 5; Conservative 0; Mismatches 0; GTGCTCCATTGATGC 15	Unknown. Unknown. Unclassified. 1 (bases 1 to 15) 1 (bases 1 to 15) Kasid, U., Gokhale, P., Dritschilo, A. and Rahman, A. Liposomes containing oligonucleotides Patent: US 612695-A 1 03-OCT-2000; Location/Qualifiers 115 - /organism="unknown" /mol_type="unassigned DNA" tch 100.0%; Score 15; DB 6; Length 15;	ARI10775 15 bp DNA linear PAT 14-Sequence 1 from patent US 6126965. ARI10775 ARI10775.1 GI:12827623
14-FEB-2001	, o		PAT 14-FEB-2001

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KEYWORDS
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Semple, S.C., Klimuk, S.K., Harasym, T., Hope, M.J.,
Cullis, P., Scherrer, P. and Debeyer, D.
Charged therapeutic agents encapsulated in lipid
containing four lipid components
Patent: US 6287591-A 15 11-SEP-2001;
                                                                                                                                                                                                 synthetic construct synthetic construct
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CQ789712
CQ789712.1 GI:45823264
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Location/Qualifiers
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                                                         /organism="synthetic construct"
/mol_type="unassigned DNA"
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Sequence 1
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1 (bases 1) to 15)
Kasid,U., Gokhale,P., Zhang,C., Dritschilo,A. and Rahman,A.
Cationic liposomal delivery system and therapeutic use thereof
Patent: US 6559129-A 3 06-MAY-2003;
Location/Qualifiers
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Kasid, U., Gokhale, P., Zhang, C., Dri
Cationic liposomal delivery system a
Patent: US 6559129-A 1 06-MAY-2003;
Location/Qualifiers
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Sequence 25 from Patent WO03094828.
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Sequence 25 from Patent
AX957646
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                   Tam,Y.K., Semple,S., Klimuk,S. and Chikh,G. Cancer vaccines and methods of using the same Patent: WO 03094828-A 25 20-NOV-2003;
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                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                      Homo
                                                                                                                                     Homo sapiens (human)
                                                                                                                                                                      AX957740.1
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Methylated immunostimulatory oligonucleotides
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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100.0%; Pred. No. 1.2e+03;
tive 0; Mismatches 0;
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Patent: JP 2002501511-A 15 15-JAN-2002;
INEX PHARMACEUTICALS CORP
PN JP 2002501511-A/15
PD 15-JAN-2002
PF 14-MAY-1998 JP 1998548646
PF 14-MAY-1998 JP 1998548646
PF 15-BAN C SEMPLE, SANDRA K KLIMUK, TROY HARASYM, MICHAEL J HOPE, PI STEVEN M ANSELL,
PI PIETER CULLIS, PETER SCHERRER, DAN SUITE DEBEYER PC A61K9/00
CC Strandedness: Single;
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Chlamydia sp.

Chlamydia sp.

Chlamydia sp.

Bacteria; Chlamydiae; Chlamydiales; Chlamydiaceae; Chlamydia.

1 (bases 1 to 15)

Semple, S.C., Klimuk, S.K., Harasym, T., Hope, M.J., Ansel, S.M.,

Cullis, P., Scherrer, P. and Debeyer, D.S.

High efficiency encapsulation of charged therapeutic agents in the desired residence.
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High efficiency encapsulation of charged therapeutic agents
lipid vesicles.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AR073978
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Monia, B.P. and Boggs, R.T.
Antisense oligonucleotide modulation of raf
Patent: US 5952229-A 47 14-SEP-1999;
Location/Qualifiers
1 GTGCTCCATTGATGC 15
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Sequence 49 from patent US
AR216002
AR216002.1 GI:23314290
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Patent: US 6410518-A 49 25-JUN-2002;
Location/Qualifiers
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/db_xref="taxon:35827"
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100.0%; Pred. No. 1.1e+03;
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Kasid, U., Gokhale, P., Dritschilo, A. and
Liposomes containing oligonuclectides
Patent: US 6126965-A 2 03-OCT-2000;
Location/Qualifiers
                                          1 (bases 1 to 25)
Kasid, U., Gokhale, P., Zhang, C., Dritschilo, A. and Rahma Cationic liposomal delivery system and therapeutic use Patent: US 6559129-A 2 06-MAY-2003;
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Aaa73535 Mouse and
Aad44760 Mouse and
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Ade90171 Human c-r
Ade30025 Antisense
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Adf41926 Methylate
Adf4170154 Oligonucl
Adf8950 Anti c-ra
Aat27527 Mouse/rat
Aaz11557 Mouse and
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Adf18639 Human raf	ADF18639	10	2977	100.0	15	45	a
Aad44819 Human raf	AAD44819	σ	2977	100.0	15	44	a
Aai68698 Human c-r	AAI68698	6	2977	100.0	15	43	a
Abk72300 Lymphona	ABK72300	σ	2977	100.0	15	42	a
Abl57050 Human pro	ABL57050	6	2977	100.0	15	41	a
Abl69999 Pancreas	ABL69999	σ	2977	100.0	15	40	n
Aaf75126 Human c-R	AAF75126	σ	2977	100.0	15	39	ဂ
Aaa73552 Polynucle	AAA73552	w	2977	100.0	15	38	ი
Aaa48654 Human c-r	AAA48654	w	2977	100.0	15	37	a
Aax78137 Human c-R	AAX78137	N	2977	100.0	15	36	a
Aax99340 Human c-r	AAX99340	N	2977	100.0	15	3 5	a
Aav20439 Human c-r	AAV20439	N	2977	100.0	15	34	ဂ
Aat61894 Human raf	AAT61894	N	2977	100.0	15	ω ω	n
Aat30085 Human Raf	AAT30085	N	2977	100.0	15	32	a
Adl16251 Raf cDNA.	ADL16251	12	2975	100.0	15	31	a
Aad64080 DNA #1 re	AAD64080	10	2975	100.0	15	30	a
Adb52556 Primary r	ADB52556	10	2524	100.0	15	29	a
Adb58074 Toxicity-	ADB58074	10	2524	100.0	15	28	a
Adf57946 Human pol	ADF57946	10	1038	100.0	15	27	ဂ
Adr60510 Cotton cD	ADR60510	13	968	100.0	15	26	
Ach77604 Human gen	ACH77604	12	597	100.0	15	25	
Ach17231 Human adu	ACH17231	φ	478	100.0	15	24	ი
Abq97799 Mouse ES	ABQ97799	σ	165	100.0	15	23	a
Adb97457 Oligo use	ADB97457	10	25	100.0	15	22	
Acd42120 Antisense	ACD42120	10	20	100.0	15	21	

RESULT 1 AAV54043 AAV54043 standard; DNA; 15 BP 02-DEC-1998 (first entry) AAV54043;

ALIGNMENTS

Human antisense c-raf-1 oligodeoxyribonucleotide.

Human; antisense; c-raf-1; oligodeoxyribonucleotide; ODN/oligo; tumour tissue; cancer; radiation therapy; radiosensitise; antisense; liposome carrier system; ss.

modified_base Homo sapiens Location/Qualifiers *tag=

modified_base WO9843095-A1 /note= /*tag= /note= "C-terminal base is phosphothioated"

"N-terminal base is phosphothioated"

21-MAR-1997; 24-OCT-1997; 01-OCT-1998. 19-MAR-1998; 97US-0041192P 97US-00957327 98WO-US005303

Kasid U, Gokhale P, Dritschilo A, Rahman

(GEOU) UNIV GEORGETOWN.

WPI; 1998-532155/45.

New cationic liposome composition containing raf oligodeoxynucleotide can be used to directly target tumour tissue and is useful in the radiation therapy of cancers.

Claim

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25pp;

English

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RESULT 2
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                The present sequence represents an antisense oligonucleotide directed against the human c-raf-1 protein kinase gene. The oligonucleotide can have either phosphorothioate or phosphodiester bonds. The oligonucleotide is lipid-encapsulated using the method of the invention. A composition comprising lipid-encapsulated particles of a therapeutic agent, e.g. antisense oligonucleotides, is prepared by mixing at least 2 lipids with buffered aqueous solution of charged therapeutic agent to form an intermediate mixture of lipid-encapsulated particles, and changing the pH of the mixture to neutralise at least some of the external surface
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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                                                                                                                                                                                                                                                                               98pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "phosphorothioate
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Τ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.1e+02;
thes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
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                                                                                                                                                                                                                                                                                                                                                     c agent - useful,
or treating dise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cullis
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human This sequence Example 5;

represents an antisense oligonucleotide protein kinase as its target gene. The

s sequence which oligonucleotide

has is

A method

for deliver comprising

delivering

ring antisense oligonucleotides steric barrier lipids.

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cells

using

lipid

Page 57;

99pp; English

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RESULT 3
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ID AAZS
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            charges on the particles. One lipid has a (de)protonatable group with Ka such that the lipid is charged at a first pH but neutral at a second pH (particularly near physiological pH) and the buffer maintains this lipid in the charged form (i.e. cationic when the therapeutic agent is anionic in the buffer, or vice versa). The second lipid prevents particle aggregation during formation of the lipid-therapeutic agent particles. The composition is used to introduce therapeutic agents into cells, in vivo or in vitro, particularly to treat or prevent diseases associated with aberrant gene expression in mammals, specifically tumours, inflammation or infection
                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense oligonucleotide; phosphorothioate; inflammatory disease; tumour; gene therapy; aberrant gene expression; treatment; infectious disease; protein kinase C alpha; c-raf-1 protein kinase
                                                                                     WPI; 2000-225058/20.
                                                                                                          Scherrer
                                                                                                                    Klimuk
                                                                                                                                                                                                                                                                13-MAY-1999;
                                                                                                                                                                                                                                                                                     14-NOV-1999.
                                                                                                                                                                                                                                                                                                          CA2271582-A1
                                                                                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ98661;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ98661 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 15 BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                              (CULL/)
                                                                                                                                                                                                                                           14-MAY-1998;
                                                                                                                                          (SEMP/)
                                                                                                                                                                                     (ANSE/)
                                                                                                                                                                                                 (HOPE/)
                                                                                                                                                                                                                    (KLIM/)
                                                                                                   SK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                c-raf-1
                                                                                                                                        HOPE M J.

ANSELL S M.

CULLIS P R.

MOK W W K.

SCHERRER P.

SEMPLE S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Н
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                                                                                                         Harasym
Semple
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                PK
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/note= "Optionally
                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                               linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapeutic antisense oligonucleotide
                                                                                                        SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                    Hope
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Pred. No. 1.1e+02;
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                                                                                                                     Ansell
                                                                                                                                                                                                                                                                                                                                         phosphorothicate
                                                                                                                     SM,
                                                                                                                     Cullis
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                                                                                                                                                                                                                                                                                                                                          internucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                    protein kinase;
                                                                                                                    PR,
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RESULT 4
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tumours. The composition comprises lipid-therapeutic agent particles comprising a lipid portion and a charged therapeutic agent which is encapsulated in the lipid portion. The lipid portion comprises a first lipid component selected from lipids containing a protonatable or deprotonatable (preferably protonatable) group that has a pKa such that the lipid is in charged form at a first pH and in neutral form at a
                                                                                                                                                                                                                     Semple SC,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      used in a method for delivering lipid encapsulated therapeutic agents (i.e antisense oligonucleotides) to mammals. The lipid capsule comprises steric barrier lipids that prevent particle aggregation during lipid nucleic acid formation. The method may be used for the delivery of therapeutic agents to mammalian cells. It is especially suitable for delivering nucleic acid molecules, and in particular antisense molecules which may be administered to down regulate the expression of aberrant genes. The aberrant gene may be ICAM-1, c-myo, ras, raf, erb-B-2, PKC-alpha, IGF-1R, EGFR, VEGF and/or VEG-R-1. The method may be used for the treatment of tumours, inflammatory diseases and/or infectious
                                                                                                                                           Composition useful for treatment of e.g. tumors con
                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                           14-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treatment; tumour; lipid-therapeutic agent particle; sphingomyelin; distearcylphosphatidylcholine; palmitcylchecyl phophatidylcholine; DSPC; pOPC; 1,2-diolecyl-sn-3-phosphoethanolamine; cholesterol; SM; DOPE; inflammation; c-raf-1 protein kinase gene; human; infectious disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD22797 standard; DNA; 15 BP
                                                                                                                                                                                                                                                                                               14-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human c-raf-1 protein kinase antisense oligonucleotide, ATG-AS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                       The invention relates to a composition useful for treatment of e.g.
                                                                                                                   Disclosure; Col 15-16; 48pp; English.
                                                                                                                                                                                                                                                                  (INEX-) INEX PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 2
                                                                                                                                                                                                                                     Klimuk SK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                       Debeyer D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                               97US-00856374
                                                                                                                                                                                                                                                                                                                            98US-00078954.
                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                  base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                 "Optionally phosphorothicate backbone"
                                                                                                                                                                                                                                     Harasym T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Other;
                                                                                                                                                                                                                                      Ansell SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 15
                                                                                                                                                              comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       infectious disease; ss
                                                                                                                                                agent
   pKa such that form at a
                                                                                                                                                              particles
                                                                                                                                                                                                                                      Cullis
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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                component is further selected such that the charged form is cationic when the therapeutic agent is anionic and vice versa; the second lipid component is selected from lipids that prevent particle aggregation during lipid-therapeutic agent particles formation and which exchange out the lipid particle at a rate greater than PEG-CerC20; third lipid component is a neutral lipid selected from distearcylphosphatidylcholine (CDC), palmitcylolecyl phosphatidylcholine (PDC), palmitcylolecyl phosphatidylcholine (PDC), palmitcylolecyl phosphatidylcholine (PDC), palmitcylolecyl phosphatidylcholine (PDC), component which is cholesterol. Compositions of the invention are used for treatment or prevention of a disease caused by aberrant expression of a gene preferably ICAM-1 (Intracellular adhesion molecule-1), c-myc, c-myb, ras, raf, carb-B-2, PKC-alpha (phosphokinase C-alpha) IGF-IR (insulin growth factor 1-receptor), bcl-2, EGFR (epidermal growth factor 1 receptor) and amammal or by inflammations such as tumour or an infectious constant of the present sequence is an antisense oligonucleotide targetted to human c-raf-1 protein kinase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                      Improved mucosal adjuvant useful in the preparation of vaccine stimulating an immune response comprises a lipid-nucleic acid containing a nucleic acid component encapsulated by a lipid.
                                                                                                                                                                                                     07-NOV-2001; 2001US-0337522P
10-MAY-2002; 2002US-0379343P
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lipid nucleic acid;
C-Raf-s; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide ODN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACC58517
                           Disclosure;
                                                                                                                                                                                                                                               07-NOV-2002; 2002WO-CA001717
                                                                                                                                                                                                                                                                                15-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                  WPI; 2003-493235/46.
                                                                                                                                                                                                                                                                                                            WO2003039595-A2
                                                                                                                                                                          (INEX-) INEX PHARM CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA; 15
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                                                                                                                                            Klimuk S,
                          Page 21; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A; 4 C; 4 G; 5 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LNA; mucosal; vaccine; immunostimulant; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #25 (hC-Raf1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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Pred. No.
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The present sequence

is that of oligodeoxynucleotide ODN #25 (hC-Raf-1)

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RESULT 6
ADA24233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 for human C-Raf-s. It is an example of an ODN that can be used in lipid-nucleic acid (LNA) formulations of the invention comprising a lipid component and a nucleic acid component. The invention is based on the discovery that such LNA formulations associated with a target antigen stimulate enhanced mucosal immune responses, especially IgA production, directed to that target antigen in vivo as compared to the target antigen alone or mixed with free or unencapsulated forms of the ODN. Claimed improved mucosal vaccines comprise an LNA formulation with at least one antigen, the LNA formulation component the INA formulation with at least one encapsulates the nucleic acid component, with the lipid and nucleic acid components acting synergistically to stimulate antigen-specific IgG production in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapeutic oligonucleotide; double-stranded RNA; dsRNA; mobile protein; cytostatic; immunosuppressive; virucide; anti-HIV; antibacterial; cardiant; hyperproliferation; cancer; haematological; metastatic;
                                                                                Treating diseases with oligonucleotides or interfering RNA, useful e.g. for cancer or autoimmune diseases, covalently coupled to mobile proteins, in vivo or in vitro.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   autoimmune disease; infection; endocrine; neural; cardiovascular; pulmonary; reproductive system disorder; endocytosis; metabolic process; murine; intracellular adhesion molecule 1; ICAM-1;
                                                    Claim 128; Page 12; 42pp;
                                                                                                                                                                                                                                                              13-FEB-2003; 2003WO-US004323.
                                                                                                                                                                                                                                                                                             21-AUG-2003
                                                                                                                                                                                                                                                                                                                         WO2003069306-A2
                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human c-raf-1 protein kinase antisense oligonucleotide SEQ ID NO:16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15
                       The present
                                                                                                                                           WPI; 2003-646491/61.
                                                                                                                                                                         Xie D;
                                                                                                                                                                                                                                  13-FEB-2002; 2002US-0356053P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense oligonucleotide; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-NOV-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           production
                                                                                                                                                                                                      (MEDB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                      MEDBRIDGE INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTGCTCCATTGATGC 15
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         invention ng: (a) a t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 A; 4 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                            /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                     note=
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                        describes
                                                                                                                                                                                                                                                                                                                                                   "optionally phosphorothicate backbone"
                                                    English.
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Pred. No.
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                       a method
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      a method for treating oligonucleotide (TON)
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disease by
or double-stranded
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administering:

therapeutic

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Chemosensitization

itization of lymphoma or

tumor tissue, useful for treating cancer, myeloma, comprises administering a chemotl

chemotherapeutic

e.g.

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RESULT 7
ADB97458/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC RNA (dsRNA) that includes a reactive group (RG) that can react with a combile protein (MP) to form a covalent conjugated of TON/dsRNA and MP; or CC (b) TON or dsRNA already conjugated to MP through a covalent bond. Also described: (1) TON of 15-30 bases that includes (i) a part that binds to target RNA or DNA and (ii) RG; (2) TON of 15-30 bases that includes a CC part that binds to target RNA or DNA and is conjugated to MP through a covalent link; (3) dsRNA that is conjugated to MP through a covalent link. TON have cytostatic, immunosuppressive, virucide, anti-HIV, antibacterial and cardiant CC immunosuppressive, virucide, anti-HIV, antibacterial and cardiant CC immunosuppressive, virucide, anti-HIV, antibacterial infection of metastatic spread); autoimmune diseases; viral or bacterial infections; cendocrine, neural, cardiovascular, pulmonary or reproductive system CC disorders. Also where TON or dsRNA are labelled, they can be used for disorders. Also where TON or dsRNA are labelled, they can be used for CTON/dsRNA have better cell entry (via endocytosis or other parts of the mobile protein metabolic process) and longer therapeutic life, increased from hours to weeks (the result of increased resistance to nuclease), cwithout loss of affinity for the target. In many cases immune response to CTON/dsRNA is also reduced, as is non-specific binding to endogenous colligonucleotide, which is a specifically claimed TON from the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                            dimyristoyl trimethyl ammonium propane; DMTAP; phosphatidylcholine; PC; cholesterol; cancer; leukaemia; lymphoma; myeloma; carcinoma; sarcoma; combination therapy; pre-cancerous lesion; chemotherapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                               antisense; ATG-S; raf ODN; chemosensitisation; tumour tissue; chemotherapeutic agent; cationic liposome; cationic lipid; phosphatidylcholine; cholesterol; liposome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sense (ATG-S) raf ODN oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB97458 standard; DNA; 15
                                                  WPI; 2003-689738/65.
                                                                                    Rahman
                                                                                                     Kasid U,
                                                                                                                                                                                                                          14-FEB-2003; 2003WO-US004681.
                                                                                                                                                                                                                                                            28-AUG-2003
                                                                                                                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB97458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                        15-FEB-2002; 2002US-00075994
                                                                                                                                                                                                                                                                                             WO2003070221-A1
                                                                                                                                                                                                                                                                                                                                                                                                                    dimethyldioctadecyl ammonium bromide;
                                                                                                                                     (GEOU ) UNIV GEORGETY (NEOP-) NEOPHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GTGCTCCATTGATGC 15
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                                                                                                                                                         UNIV GEORGETOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                     Gokhale P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 C; 4 G;
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                                                                                                   Pei J,
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Pred. No. 1.1e+02;
                                                                                                     Mewani
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                                                                                                     ₽,
                                                                                                     Ahmad
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                                                                                                     Drischilo A;
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Example 1; Page 18; 77pp; English

agent and cationic liposomes containing oligonucleotides

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RESULT 8
ADB97456
ID ADB9
XX ADB9
XX ADB9
XX ADB9
XX ADB9
XX ADB9
XX ANTi
XX ANTi
XX ANTi
XX ANTi
XX Chem
KW Choll
XX Choll
XX Comb
XX Comb
XX WO20
XX WO20
XX WO20
XX WO21
XX WO21
XX WO21
XX WO21
XX WO21
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XX GEC
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XX GE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chemotherapeutic agent; cationic liposome; cationic lipid; phosphatidylcholine; cholesterol; liposome; dimethyldioctadecyl ammonium bromide; DDAB; dimyristovl + rime+k-i ~---
                                                         Chemosensitization of tumor tissue, useful for treating leukemia, lymphoma or myeloma, comprises administering a agent and cationic liposomes containing oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dimyristoyl trimethyl ammonium propane; DMTAP; phosphatidylcholine; PC; cholesterol; cancer; leukaemia; lymphoma; myeloma; carcinoma; sarcoma;
                                                                                                                                                                                    WPI; 2003-689738/65.
                                                                                                                                                                                                                                                                          Kasid U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-FEB-2003; 2003WO-US004681.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-AUG-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003070221-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            combination therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense (ATG-AS) raf ODN oligodeoxyribonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15
                                                                                                                                                                                                                                          Rahman A;
                                                                                                                                                                                                                                                                                                                                                            (GEOU ) UNIV GEORGETOWN.
                                                                                                                                                                                                                                                                                                                                                                                                                     15-FEB-2002; 2002US-00075994
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                                                                                                                                                                                                                                                                                                                                (NEOP-) NEOPHARM INC.
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                                                                                                                                                                                                                                                                      Gokhale P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pre-cancerous
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                                                                                                                                                                                                                                                                      Mewani R,
                                                                                           comprises administering a chemotherapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lesion; chemotherapy; ss.
                                                            oligonucleotides
                                                                                                                                                                                                                                                                      Ahmad I,
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Example 1; Page 18; 77pp; English

Disclosure;

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25; 138pp; English

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RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC The invention relates to a novel method for the chemosensitisation of CC tumour tissue, comprising administering a chemotherapeutic agent and a CC composition comprising cationic lipsosmes consisting of cationic lipid, CC phosphatidylcholine and cholesterol, where oligonuclectide(s) are CC encapsulated within the liposome. The invention further relates to a composition comprising liposomes consisting essentially of a cationic CC lipid like dimethyldioctadecyl ammonium bromide (DDAB) or dimyristoyl CC trimethyl ammonium propane (DMTAP), phosphatidylcholine (PC), where CC cholesterol, and containing the sequence 5'- GTGCTCCATTGATGC -3', where CC conjy the terminal sequences are phosphorothioated. The method is useful CC lymphoma, myeloma, carcinoma or sarcoma. The combination therapy may be CC used for any stage of cancer ranging from pre-cancerous lesions to cancer of advanced stages. This polynucleotide sequence represents the antisense CC (ATG-AS) raf ODN oligodeoxyribonucleotide, a cationic liposome of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 15
                                          New pathogen vaccine having a lipid-nucleic acid formulation combination with at least one microbial antigen, useful for enhanced responses against bacterial, viral and parasitic inf
                                                                                                                                                                                       10-MAY-2002; 2002US-0379343P
07-NOV-2002; 2002US-00290545
12-MAR-2003; 2003US-0454298P
                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                   WPI; 2003-903935/82
                                                                                                                                                                                                                                                                                  20-NOV-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004
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                                                                                                                                                                                                                                                                                                                WO2003094829-A2
                                                                                                                                                             (INEX-) INEX PHARM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunostimulant; vaccine; lipid-nucleic acid; phosphorothioate; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunostimulant ODN25, component of lipid-nucleic acid vaccine
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                                                                                                                              chikh G,
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                           "OTHER= optional phosphorothioate nucleotides"
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Matches 15
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                                                                                                                                                                                                                                                                                                    Semple SC,
Scherrer P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; lipid-encapsulated therapeutic agent particle; aberrant gene expression; intercellular adhesion molecule; ICAM-1; c-myc; c-myb; ras; raf; erb-B-2; protein kinase C; PKC-alpha; insulin-like growth factor; IGF-IR; epidermal growth factor receptor; EGFR; vascular endothelial growth factor; VEGF; VEGF-R-1; tumour;
                                                                                                            agent partic.
                                                                                                               Preparation of a composition comprising lipid-encapsulated therapeutic agent particles, useful for introducing a nucleic acid into a cell and for treating diseases characterized by aberrant gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SEMP/)
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HOPE M J.
ANSELL S M.
CULLIS P.
SCHERRER P.
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Debeyer D
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98US-00078954.
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100.0%; Pr
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Pred. No.
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ches 0;
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Disclosure; SEQ

ID NO 15;

lipid-nucleic acid (INA) formulation in combination with at least one tumour-associated antigen that is mixed with or associated with the IN formulation comprising a lipid component having at least one cationic lipid, and a nucleic acid component comprising at least one cationic oligonucleography.

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Best Local S
Matches 15
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                                                                                                                      New cancer vaccine having a lipid-nucleic acid formulation in combination with at least one tumor-associated antigen, useful for stimulating enhanced responses against tumor-associated antigens and for inhibiting
                                                                                                                                                                                                                                              10-MAY-2002; 2002US-0379343P
07-NOV-2002; 2002US-00290545
04-APR-2003; 2003US-0460646P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide ODN 25 (hC-Raf-1) SEQ ID NO:25.
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                                                                                                                   enhanced responses against
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer; vaccine; lipid-nucleic acid; LNA; tumour-associated antigen;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                response; cytostatic; gene
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Pred. No.
                                                                                  English
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hes 0;
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                                                                                                                                                                                                                                                                                              The present invention relates to reducing the platelet count in a patic comprises preparing a formulation of an oligonuclectide with an agent that enhances penetration of the oligonuclectide into cells, and administering the formulation to a patient having an elevated platelet count. The oligonuclectide is useful for preparing a medicament for reducing the platelet count in a patient, particularly for treating the present sequence represents an oligonuclectide of the plate 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Reducing the platelet count in a patient, useful for treating thrombocytosis, comprises administering antisense oligonucleotides inhibiting raf-1 gene with an agent that enhances penetration of the oligonucleotide into cells.
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                                                                                                                                                                                                                     Sequence 15
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                                                      GTGCTCCATTGATGC 15
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                                                                                                             Conservative
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; Mismatches 0;
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Pred. No. 1.1e+02;
; Mismatches 0;
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                                                                                                                                                              Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                  CC formulation for stimulating an immune response in an animal, comprising a CC lipid component and a nucleic acid component which is a methylated cullipid component and a nucleic acid component which is a methylated cc nucleic acid (LNA) formulation; (2) a vaccine comprising a lipid component in combination with at least one target antigen; (3) cc stimulation in combination with at least one target antigen; (3) cc stimulating an enhanced host immune response to antigenic stimulation, cc comprising administering to the host the LNA formulation; (4) stimulating comprising contacting at least one cc dendritic cells in vivo, comprising contacting at least one cc dendritic cell with the lipid-methylated nucleic acid formulation to a cc stimulation to antigen presenting cells, comprising the administration of the LNA formulation associated with a target antigen. The lipid-cc methylation nucleic acid formulation has immunostimulant, cytostatic, and in gene therapy. The formulation and methods are useful in crimina trains a transfer and enhanced are useful in the property of the complete of the complete contacting and methods are useful in the component of the complete contacting and can be useful in the component of the contacting are useful in the contacting and methods are useful in the contacting and contacting are useful.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-MAY-2002; 2002US-0379343P
07-NOV-2002; 2002US-00290545
04-APR-2003; 2003US-0460646P
                                                                                                                                                                                                                                                                                                     stimulating a host's immune response to antigenic stimulation, or activating and/or expanding dendritic cell populations in response antigenic stimulation. They may be used for treating cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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arthritis; immunodeficiency disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     lipid-methylated nucleic acid formulation; immune response; lipid-nucleic acid; vaccine; immunostimulant; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF42926 standard; DNA; 15
                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-MAY-2003; 2003WO-CA000678.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present
                                                                                                                                                                                                     sequence represents a methylated immunostimulatory in the exemplification of the present invention.
                                                                                                                                                                                                                                                                         inflammation, arthritis or immunodeficiency disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INEX-) INEX PHARM CORP.
                                   Similarity
                                                                                                                                           15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunostimulatory oligonucleotide;
                                                                                                                                        вP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention describes a lipid-methylated nucleic acid
       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                        2 A;
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                                                                                                                                        4 C; 4 G;
                                   100.0%;
                                                                100.0%;
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   Score 15; DB 12;
Pred. No. 1.1e+02;
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                                                                                                                                        Ή,
                                                                                                                                           0 U;
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                                                                       Length 15
                                                                                                                                                                                                                                        ers. The present oligonucleotide given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ
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RESULT 14
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is that of an antisense oligonucleotide to raf. The invention relates to novel gemcitabine compositions and their use in treating proliferative diseases such as cancer, particularly in mammals, especially in humans. The compositions include liposome-entrapped gemcitabine. The cancer is especially lymphoma, ovarian cancer, breast cancer, pancreatic cancer, lung cancer or colon cancer. The liposomal gemcitabine compositions can be used in conjunction with secondary therapeutic agents including antineoplastic, antifungal and antibiotic agents as well as antisense oligonucleotides, especially an antisense oligonucleotide to raf (claimed).
                                                                                                                                          C-raf-1; liposomal; antineoplastic; cytostatic; cancer; antisense; ss.
                                                                                                                                                                                                                                                                       18-NOV-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 1; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treatment of cellular proliferative disease e.g. cancer involves administration of a composition comprising liposomal gemcitabine and negatively charged phospholipid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-AUG-2003; 2003WO-US025293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide antisense to raf.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAY-2004
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                                                                                                                                                                                                  Anti c-raf-1 oligonucleotide
                                                                                                                                                                                                                                                                                                                                   ADR88950;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 15 BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
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                WO2004071466-A2
                                                                                                                                                                                                                                                                                                                                                                                                 ADR88950 standard;
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Search completed: June 21, Job time : 438 secs

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method involves; dissolving a lipid fraction in a water-miscible organic solvent; and mixing the water-miscible organic solvent solution comprising the lipid fraction with an aqueous solution under conditions to form a bulk liposomal composition. The method further involves adding at least one active principal to the water-miscible organic solvent prior to the addition of the lipid fraction. The method further involves adding composition of the lipid fraction, or to the aqueous solution prior, composition to obtain a size-reducing the bulk liposomal composition, freeing the liposomal composition, freeing the composition, sterile-filtering the liposomal composition, freeing the liposomal composition, sterile-filtering the liposomal composition and sterile corganic solvent by the liposomal preparation. Step (b) involves adding the water-miscible organic solvent prior, the aqueous solution while mixing and mixing of solution following addition of water-miscible solvent comprising the lipid fraction to the aqueous solution while mixing and mixing composition to the aqueous solution while cooling. The active principal comprises at least one antineoplastic or antifungal agent (preferably taxane, camptothecin or their derivatives, especially compositions, need not composition is used for the treatment of disease e.g. cancer. The composition is used for the treatment of compositions pred not composition on a commercial scale. The present sequence represents an antisense oligonucleotide specific for c-raf-1, that can be used as an active principal.
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Matches
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                                                                                                                                                             Sequence 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Preparation of liposomal composition used for treating e.g. cancer involves dissolving lipid fraction in water miscible organic solvent and mixing solvent solution with aqueous solution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bhamidipati S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-FEB-2003; 2003US-0446895P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to the preparation of a liposomal composition. The
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                                                                                                   Local
                                                                               . Similarity
15; Conserv
                                      1 GTGCTCCATTGATGC 15
GTGCTCCATTGATGC
                                                                                                                                                               BP; 2 A; 4 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                 Conservative
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15
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Pred. No. 1.1e+02;
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